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April 2, 2004, 10:04:38; Search time 31.3333 Seconds (without alignments) 151.046 Million cell updates/sec
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GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
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                                                                                                                                                                                                                                                          1017041 seqs, 315518202 residues
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Maximum Match 100%
Listing first 45 summaries
                                                            OM protein - protein search, using sw model
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Gapop 10.0 , Gapext 0.5
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_vertebrate:*
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sp_human:*
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sp_bacteria:*
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Maximum DB seq length: 15
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112
113
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144
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                                                                                                                                                                         Perfect score:
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pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

	Description	Q8cja8 mus musculu	Q8cja9 musculu	Q945c1 crypthecodi	O98696 hordeum vul	Q13865 homo sapien	Q9twv0 anthopleura	Q8jv78 polyomaviru	O88176 mus musculu		Ogeuz3 escherichia	018890 ateles belz	Q9g3c0 calliphora	O14182 homo sapien	Q98905 glycine max	рошоц	Q9ucc2 homo saplen
	ជា	OBCJA8	OBCJA9	0945C1	969860	013865	Q9TWV0	Q8JV78	088176	Q8GTG5	O9EUZ3	018890	096300	014182	098905	80NI60	Q9UCC2
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ALIGNMENTS

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MEDLINE=22257759; PubMed=12370778;
Tran P., Leclerc D., Chan M., Pai A., Hiou-Tim F., Wu Q., Goyette P.,
Artigas C., Miloson R.,
"Multiple transcription start sites and alternative splicing in the
methylenetetrahydrofolate reductase gene result in two enzyme
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Butheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
NCBI_TaxID=10090;
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0
                                                01-MAR-2003 (TrEMBLrel. 23, Created)
01-MAR-2003 (TrEMBLrel. 23, Last sequence update)
01-MAR-2003 (TrEMBLrel. 24, Last annotation update)
Methylenetetrahydrofolate reductase short isoform (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                       Score 27; DB 11; Length 14;
pred. No. 8.7e+02;
1; Mismatches 5; Indels
                                                                                                                                                                                                                                                                                                                        isoforms.";
Mamm. Genome 13:483-492(2002).
EMBL; AF404271; AAN40873.1; -.
MGD; MGI:106639; Mthfr.
14 14 14
SEQÜENCE 14 AA; 1472 MW; 28DD341AC1695CB8 CRC64;
                        14 AA.
                        PRT;
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Best Local Similarity 45.5%;
Matches 5; Conservative
                         PRELIMINARY;
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                                                                                                                                  Mus musculus (Mouse).
                                                                                                                                                                                                            SEQUENCE FROM N.A.
                                             QBCJAB;
                           Q8CJA8
RESULT 1
            QBCJAB
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RESULT 2 Q8CJA9

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BETA-KINESIN
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Crypthecodinium cohnii (Dinoflagellate).
Eukaryota; Alveolata; Dinophyceae; Gonyaulacales; Crypthecodiniaceae;
Crypthecodinium.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                STRAIN=129/Sv;
MEDLINE=22257759; PubMed=12370778;
Tran P., Leclerc D., Chan M., Pai A., Hiou-Tim F., Wu Q., Goyette P., Artigas C., Milos R., Rozen R.;
"Multiple transcription start sites and alternative splicing in the methylenetetrahydrofolate reductase gene result in two enzyme
                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                         Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Rodentia, Sciurognathi, Muridae, Murinae, Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0
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                       01-MAR-2003 (TrEMBLrel. 23, Created)
01-MAR-2003 (TrEMBLrel. 23, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
Methylenetetrahydrofolate reductase short isoform (Fragment)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 26; DB 10; Length 13;
Pred. No. 1.2e+03;
1; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                         34.2%; Score 27; DB 11; Length 15; 45.5%; Pred. No. 9.3e+02; tive 1; Mismatches 5; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A. MEDLINE=2142816; MEDLINE=21428164; PubMed=11545436; Guillebault D., Derelle E., Bhaud Y., Moreau H.; "Role of nuclear WW domains and proline-rich proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              13 AA; 1225 MW; 322FA05EE70CB327 CRC64;
                                                                                                                                                                                                                                                                     SEQUENCE 15 AA; 1559 MW; 8D78DD341AC1695C CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                         (TrEMBLrel. 19, Created)
(TrEMBLrel. 19, Last sequence update)
(TrEMBLrel. 19, Last annotation update)
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01-MAY-1999 (TrEMBLrel. 10, Created)
01-MAY-1999 (TrEMBLrel. 10, Last sequence update)
                                                                                                                                                                                                                                                                                                                                                                                                                     13 AA.
     15 AA.
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      PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      dinoflagellate transcription.";
Protist 152:127-138(2001).
EMBL; AF417570; AAL15908.1; -.
                                                                                                                                                                                                                                       EMBL; AF404270; AAN40872.1; -. MGD; MGI:106639; Mthfr.
                                                                                                                                                                                                                             Mamm. Genome 13:483-492(2002).
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55.6%;
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Matches 5; Conservative
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                                                                                                                                                                                                                                                                                                                    5; Conservative
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        PRELIMINARY;
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                                                                                    Mus musculus (Mouse)
                                                                                                                                                                                                                                                                                                          Best Local Similarity
                                                                                                                                          SEQUENCE FROM N.A.
                                                                                                                   NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NCBI_TaxID=2866;
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QBCJA9;
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098696
ID 0986
AC 0986
DT 01-M
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ENBL; X69658; CAA43349.1; ---
PIR; S43170; S43170.
                                                                                                                                                                                                         STRAIN-Hassan, TISSUE-Leaf,
MEDINE-2013023, PubMed=10666448,
del Campo E., Sabater B., Martin M.,
"Transcripts of the ndhH-D operon of barley plastids: possible role of
unedited site III in splicing of the ndhA intron.";
Nucletc Acids Res. 28:1092-1098(2000).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                               Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; Pooideae;
Triticeae; Hordeum.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Bukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE 14 AA; 1521 MW; 361CF5911A68F8B8 CRC64;
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Last annotation update)
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01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
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55.6%; Pred. No. 2.3e+03;
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Pred. No. 1.9e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        31.6%;
80.0%;
                             ORF321 protein (Fragment). Hordeum vulgare (Barley).
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01-JUN-2003 (TrEMBLrel.
Kinesin light chain.
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Best Local Similarity
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EMBL; AJ132862; CAC20133.1; -.
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                       0; Gaps
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDLINB=21465052; PubMed=11581397;
Bofill-Mas S., Formiga-Cruz M., Clemente-Casares P., Calafell F.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Girones R.; potential transmission of human polyomaviruses through the gastrointestinal tract after exposure to virions or viral DNA."; J. Virol. 75:10290-10299(2001).
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0
  Anthopleura elegantissima (Sea anemone).
Eukaryota; Metazoa; Cnidaria; Anthozoa; Zoantharia; Actiniaria;
Nynantheae; Actiniidae; Anthopleura.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                29.1%; Score 23; DB 12; Length lu 50.0%; Pred. No. 2.9e+03; Indels
                                                                                                                                                                                                                                                                                           29.1%; Score 23; DB 5; Length 9; 80.0%; Pred. No. 1e+06;
                                                                                                                                                                                                                                                                                                                                       1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE 10 AA; 1183 MW; 4CD6A96731A32763 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           01-0CT-2002 (TrEMBLrel. 22, Created)
01-0CT-2002 (TrEMBLrel. 22, Last sequence update)
01-0CT-2002 (TrEMBLrel. 22, Last annotation update)
                                                                                                                                                                                                                                                   9 AA; 943 MW; 2908176737686777 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      10 AA.
                                                                                                                                                                                                                                                                                                                                           0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  01-NOV-1998 (TrEMBLrel. 08, Created)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.
STRAIN=Balb-c; TISSUE=Liver;
MEDLINE=98250618; PubMed=9582442;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRT;
                                                                                                                                 MEDLINE=93126143; PubMed=1480510;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EMBL; AF303946; AAM97796.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Large T antigen (Fragment).
                                                                                                                                                                                                                         anemones.";
Peptides 13:851-857(1992).
                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity 80.04
Lac 4; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                4; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mus musculus (Mouse).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         6 FGSPRPLP 13
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                                                                                                                                                                                                                                                                                                                                                                                       9 PRPLP 13
                                                                                                                                                                                                                                                                                                                                                                                                                               3 PGPLP 7
                                                                        NCBI_TaxID=6110;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
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Būkaryota, Viridiplantae, Streptophyta, Embryophyta, Tracheophyta,
Spermatophyta, Magnoliophyta, eudicotyledons, core eudicots, asterids,
lamiids, Solanales, Solanaceae, Solanum.
Kawahigashi H., Harada Y., Asano A., Nakamura M.;
"A cis-acting regulatory element that affects the alternative splicing of a muscle-specific exon in the mouse NCAM gene.";
Blochim. Biophys. Acta 1397:305-315(1998).
EMBL, AB001873; BA31275.1; -.
MGD; MGI:97281; Ncaml.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bacteria; Proteobacteria; Gammaproteobacteria; Enterobacteriales;
Enterobacteriaceae; Escherichia.
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Cagna G., Boes A., Ulker B., Zhou A., Wanke D., Somssich I.E.
Cagna G., Boes A., Ulker B., Zhou A., Wanke D., Somssich I.E.
"Comparison of WRKY group II transcription factors from plant
Submitted (SEP-2002) to the EMBL/GenBank/DDBJ databases.
EMBL, AY157059; AAN71729.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-WAR-2003 (TrEMBLrel. 23, Created)
01-WAR-2003 (TrEMBLrel. 23, Last sequence update)
01-WAR-2003 (TrEMBLrel. 23, Last annotation update)
WRKY transcription factor IId-1 splice variant 2 (Fragment).
Lycopersicon esculentum (Tomato).
                                                                                                                                                                                                                                                                                Score 23, DB 11; Length 13; Pred. No. 3.7e+03; 1; Mismatches 0; Indels
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                                                                                                                                                                       01-MAR-2001 (TrEMBLrel. 16, Created)
01-MAR-2001 (TrEMBLrel. 16, Last sequence update)
01-MAR-2001 (TrEMBLrel. 16, Last annotation update)
Ribosome binding factor A (Fragment).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              8 AA.
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80.0%;
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Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha; Oestroidea;
Calliphoridae; Calliphora.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDLINE=98267189; PubMed=9601975; Canavez F., Moreira M.A.M., Bonvicino C.R., Parham P., Seuanez H.N.; Canavez F., Moreira M.A.M., Bonvicino C.R., Parham P., Seuanez H.N.; "Comparative gene assignment in Ateles paniscus chamek (Platyrrhini, Primates) and man: association of three separate human syntenic groups and evolutionary considerations.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Wallman J.F., Donnellan S.C.; "The utility of mitochondrial DNA sequences for the identification of forensically-important blowflies (Diptera: Calliphoridae) in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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Ateles belzebuth chamek (Chamek spider wonkey).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Platyrrhini; Cebidae; Atelinae; Ateles.
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                                                                                          Score 22; DB 2; Length 11; Pred. No. 4.6e+03;
                                                                                                                                                     3, Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EMBL; AF029165; AAB84196.1; -
GO; GO:0004295; F:trypsin activity; IEA.
GO; GO:0006295; P:proteolysis and peptidolysis; IEA.
InterPro; IPR001254; Peptidase S1.
PROSITE; PS00134; TRYPSIN_HIS; 1.
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01-MAR-2001 (TrEMBLrel. 16, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
                                     6B234CFE740879CB CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-JAN-1998 (TrEMBLrel. 05, Created)
01-JAN-1998 (TrEMBLrel. 05, Last sequence update)
01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
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                                                                                                                                                     1; Mismatches
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            11
1319 MW;
                                                                                             Query Match
Best Local Similarity 50.0%;
Matches 4; Conservative
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                                                                                                                                                                                                                                                                                  2 AKEFGRPQ 9
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11 AA;
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SEQÜENCE
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Sukaryota, Viridipantae, Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids;
eurosids I; Fabales; Fabaceae; Papilionoideae; Phaseoleae; Glycine.
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Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo.
NCBI TaxID=9606;
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                                                                                   Length 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                13 AA; 1442 MW; 25B7D365F34FC408 CRC64;
                                                     14 AA; 1720 MW; B659D9D3D460BFC6 CRC64;
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01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-JUN-2002 (TrEMBLrel. 21, Last annotation update)
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                                                                                                                                                                                                                                                                                 Last sequence update)
                                                                                  Score 22; DB 8; I
Pred. No. 5.9e+03;
                                                                           27.8%; Scc...
80.0%; Pred. No. >...
0; Mismatches
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01-NOV-1996 (TrEMBLrel. 01, Last sequent
01-UTN-2003 (TrEMBLrel. 24, Last annotat
Deoxymucleotidyltransferase (Fragment):
EMBL; AY012565; AAG42941.1; -.
GO; GO:0005739; C:mitochondrion; IEA.
Mitochondrion.
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MEDLINE=9223221; PubMed=1368037;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Hirano H., Kagawa H., Okubo K.;
Phytochemistry 31:731-735(1992).
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                                                                                                  Local Similarity 80.0 nes 4; Conservative
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                                                                                                                                                                                                                                                                                                                                              Homo sapiens (Human)
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MEDLINE=92212461; PubMed=1557127;
MEDLINE=92212461; PubMed=1557127;
Wei M.L., Cresswell P.;
"HLA-A2 molecules in an antigen-processing mutant cell contain signal sequence-derived peptides.";
Nature 356,443-446[1992).
PIR; 138246; 138246.
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01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
HLA-A2 associated peptide (Fragment).
HLA-A2 associated peptide (Fragment).
ENARAYOTA: Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
Query Match 26.6%; Score 21; DB 10; Length 10; Best Local Similarity 60.0%; Pred. No. 6.2e+03; Matches 3; Conservative 2; Mismatches 0; Indels
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SEQÜENCE 13 AA; 1355 MW; 3E111BA716EA1768 CRC64;
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4 TFGTP 8
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Q9TNQ8
DD O9TNQ
DT 01-MA
DT 01-MA-A
DT 01-MA-A
DF HOMO
OC BURART
OX NCBI
RX NCBI
RX WEDLI
RA WEDLI
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Search completed: April 2, 2004, 10:11:06 Job time : 32.3333 secs

7 GSPRPLPVV 15 | | | | | | 5 GGPRGLLAV 13

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Query Match
Best Local Similarity
Matches 7; Conserv
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3, Appli
29, Appl
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Sequence 92, Appl
Sequence 92, Appl
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                                                          April 2, 2004, 10:06:43 ; Search time 14 Seconds (without alignments) 55.314 Million cell updates/sec
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2: /cgn2_6/ptodata/2/iaa/5B_COMB.pep:*
3: /cgn2_6/ptodata/2/iaa/6A_COMB.pep:*
4: /cgn2_6/ptodata/2/iaa/6B_COMB.pep:*
5: /cgn2_6/ptodata/2/iaa/PCTUS_COMB.pep:*
6: /cgn2_6/ptodata/2/iaa/PCTUS_COMB.pep:*
GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
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US-08-602-998A-92
US-08-578-865-92
US-09-500-124-92
US-09-108-168B-3
US-08-36-168B-3
US-08-36-343A-29
US-08-476-515A-29
US-08-602-999A-255
US-09-500-124-255
US-09-500-124-255
US-09-500-124-255
US-09-500-124-86
US-08-602-999A-86
US-08-602-999A-86
US-08-602-999A-86
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US-08-602-999A-62
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US-09-500-124-62
US-07-756-250-3
US-08-602-999A-67
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US-09-500-124-67
US-08-230-047-29
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US-08-652-877-36
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                                                                                                                                                                         389414 seqs, 51625971 residues
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Maximum Match 100%
Listing first 45 summaries
                                         OM protein - protein search, using sw model
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Gapop 10.0 , Gapext 0.5
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Match Length DB
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seq length: 15
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Sequence:
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Sequence 34, Appl Sequence 16, Appl Sequence 281, Appl Sequence 281, Appl Sequence 85, Appl Sequence 101, Appl Sequence 311, Appl Sequence 315, Appl Sequence 315, Appl Sequence 114, Appl Sequence 114, Appl Sequence 114, Appl Sequence 118, Appl Sequence 378, Appl Sequence 118, Appl Sequence 118, Appl Sequence 378, Appl Sequence 118, Appl Sequence 378, Ap		SH3 BINDING
11 3 US-08-476-515A-34 11 3 US-08-476-515A-36 112 4 US-08-672-995A-281 113 3 US-08-602-999A-281 113 4 US-08-502-124-281 113 4 US-08-502-124-85 113 4 US-08-502-124-85 115 3 US-08-602-999A-355 115 4 US-09-500-124-301 115 4 US-09-500-124-35 117 4 US-09-500-124-35 118 4 US-09-500-124-35 119 4 US-09-508-188B-114 110 4 US-09-68B-188B-114 111 4 US-09-291-417D-18 112 4 US-09-291-417D-18 113 3 US-08-62-999A-35	ALIGNMENTS	ation US/08230047 Miss III, George H. shcheako, Yuri D. y, Michael C. ton: EXPRESSION CLONING OF C-SRC ton: BYPRESSION CLONING OF C-SRC ton: PROTEIN NOES: 40 NODES: 40
228 320 331 331 331 332 333 334 335 336 337 336 337 336 336 337 336 336 336		RESULT 1 US-08-230-047-30 Sequence 30, Application US/082300 Patent No. 5541109 GENERAL INFORMATION: APPLICANT: Searfoes III, Georg APPLICANT: Ivashchenko, Yuri D APPLICANT: Uspen Michael C. TITLE OF INVENTION: EXPRESSION TITLE OF INVENTION: EXPRESSION TITLE OF INVENTION: EXPRESSION CORRESEES: Rhone-Poulenc Ror STREET: 500 Arcola Road, 3C4 CITY: Collegeville STATE: PA COUNTRY: USA COMPUTER: READABLE FORM: MEDIUM TYPE: Floppy disk COMPUTER: Macintosh OPERATION SYSTEM: System 7.1 SOFTWARE: Word 5.0 (Patentify) ATTORNATION NUMBER: US/08/23 FILING DATE: 19-APR.1994 CLASSIFICATION WINBER: 32,534 REPERENCE/DOCKET UNBER: 31,534 REPERENCE/DOCKET UNBER: 32,534 REPERENCE/DOCKET UNBER: 31,534 TELEPRONE: (610) 454-3808 INFORMATION FOR SEQ ID NO: 30: SEQUENCE CHARACTERISTICS: LENGTH: 15 amino acids TYPE: amino acids TYPE: amino acids TYPE: peptide

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Gaps

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46.8%; Score 37; DB 1; Length 15; llarity 70.0%; Pred. No. 17; Conservative 0; Mismatches 3; Indels

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4 HSFGSPRPLP 13 нѕнғнркрір 10

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44.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEPHONE: (703) 413-300
TELEFAX: (703) 413-220
TELER: 248855 OPAT UR
INFORMATION FOR SEQ ID NO: 9
SEQUENCE CHARACTERISTICS:
                                                                                                  Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 62...
S; Conservative
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: C
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CORRESPONDENCE ADDRESS:
                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
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                         COUNTRY:
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STATE:
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APPLICANT: KAY, BRIAN K.
APPLICANT: SPARKS, ANDREW B.
APPLICANT: THORN, JUDITH M.
APPLICANT: QUILLIAM, LAWRENCE A.
APPLICANT: QUILLIAM, LAWRENCE A.
APPLICANT: DER, CHANNING J.
TITLE OF INVENTION: Solc SH3 BINDING PEPTIDES AND METHODS OF TITLE OF INVENTION: ISOLATING AND USING SAME
NUMBER OF SEQUENCES: 106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
ADDRESSEE: OBLON, SPIVAK, MCCLELLAND, MAIER & NEUSTADT,
ADDRESSEE: P.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              44.3%; Score 35; DB 3; Length 13; 62.5%; Pred. No. 30; cive 3; Mismatches 0; Indels
                                                                                                          APPLICANT: STARKS, Andrew B.
APPLICANT: KAY, Brian K.
APPLICANT: THORN, Judith M.
APPLICANT: OLILLIAM, Lawrence A.
APPLICANT: DER, Channing J.
APPLICANT: FOWLKES, Dana M.
APPLICANT: RIDER, James B.
TITLE OF INVENTION: ISOLATING AND USING SAME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER FRADABLE FORM:
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
COMPUTER: TBM PC COMPATION
SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATE:
APPLICATION NUMBER: US/08/602,999A
FILING DATE: 16-FBB-1996
CLASSIFICATION: 435
ATTORNEY/AGBNT INFORMATION:
NAME: MISTOCK, S. Leelle
REGISTRATION NUMBER: 18,872
REFERENCE/DOCKET NUMBER: 1101-202
TELECHOMONICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES: 467
CORRESPONDENCE ADDRESS:
ADDRESSEE: Pennie & Edmonds
STREET: 1155 Avenue of the Americas
CITY: New York
                            Sequence 92, Application US/08602999A, Patent No. 6184205, GENERAL INFORMATION:
APPLICANT: SPARKS, Andrew B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; Sequence 92, Application US/08278865; Patent No. 6303574
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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Best Local Similarity 62.5-
Local Similarity 52.5-
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; MOLECULE TYPE: peptide
US-08-602-999A-92
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RY: U.S.A.
10036-2711
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US-08-278-865-92.
           US-08-602-999A-92
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Pred. No. 30;
3; Mismatches 0; Indels
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APPLICANT: KAY, Brian K.
APPLICANT: THORN, Judith M.
APPLICANT: QUILLIAM, Lawrence A.
APPLICANT: FOWLKES, Dana M.
APPLICANT: RIDER, Channing J.
APPLICANT: RIDER, James B.
TITLE OF INVENTION: SH3 BINDING PEPTIDES AND METHODS OF TITLE OF INVENTION: 1500ATING AND USING SAME NUMBER OF SEQUENCES: 467
7: 1755 S. Jefferson Davis Highway, Suite 400
Arlington
Virginia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ZIP: 10036-2711
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PatentIn PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATE:
APPLICATION NUMBER: US/09/500,124
                                                                                                                                                                                                                    SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/278,865
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ...urkESSEE: Pennie & Edmonds
STREET: 1155 Avenue of the Americas
CITY: New York
STATE: New York
COUNTRY: U.S.
                                                                                                                                                                                                                                                                                                                                                                                                        4980-007-0
                                                                                                                                                                                                                                                                        APPLICATE:
FILING DATE:
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: Villacortex, Gilberto M.
REGISTRATION NUMBER: 34,038
REFERENCE/DOCKET NUMBER: 4980-0/
TELECOMMUNICATION INFORMATION:
""TEDHONE: (703) 4-13-3000
                                                                                                                                                                            COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          08/602,999
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APPLICANT: INFORMATION:
APPLICANT: Gastelhano, Arlindo
APPLICANT: Witter, David
TITLE OF INVENTION: CONFORMATIONALLY CONSTRAINED PEPTIDOWIMETICS AS BETA-TURN
TITLE OF INVENTION: TEMPLATES AND MODULATORS OF SH3 DOMAINS
FILE REFERENCE: 60384-8
CURRENT PELING DATE: 1998-05-28
NUMBER OF SEQ ID NOS: 5
SOFTWARE: Patentin version 3.1
SEQ ID NO 3
LENGTH: 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0
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| Sequence 29, Application US/08336343A |
| Patent No. 5677144 |
| GENERAL INFORMATION: Arel |
| APPLICANT: Ulrich, Axel |
| APPLICANT: Ulrich, Axel |
| TILLE OF INVENTION: CCK-2, A No. 5677144el Receptor Tyrosine Kinase |
| TILLE OF INVENTION: CK-2, A No. 5677144el Receptor Tyrosine Kinase |
| NUMBER OF SEQUENCES: 43 |
| NUMBER OF SEQUENCES: 43 |
| ADDRESSEE: Pennie & Edmonds |
| STREET: 1155 Avenue of the Americas |
| CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Gaps
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                                                                                                                                                                                                                                                                                                                                                                            44.3%; Score 35; DB 4; Length 13; 62.5%; Pred. No. 30; tive 3; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         43.0%; Score 34; DB 4; Length 9; 100.0%; Pred. No. 3e+05; tive 0; Mismatches 0; Indels
ATTORNEY/AGENT INFORMATION:
NAME: Mistrock, S. Lealls
REGISTRATION NUMBER: 16,872
REFERENCE/DOCKET NUMBER: 1101-202
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 790-9090
TELEPAX: (212) 869-9741/8864
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME/KEY: MISC FEATURE LOCATION: (1)...(9) OTHER INFORMATION: Synthetic Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OTHER INFORMATION: SH3 test peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 3, Application US/09086168B Patent No. 6638941
                                                                                                                                                               TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 92:
SEQUENCE CHARACTERISTICS:
LENGTH: 13 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: Artificial Sequence
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Best Local Similarity 100.0
                                                                                                                                                                                                                                                                                                                                                                                                                                      5; Conservative
                                                                                                                                                                                                                                                                                                    MOLECULE TYPE: peptide US-09-500-124-92
                                                                                                                                                                                                                                                                                       TOPOLOGY: unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     8 SPRPLPVV 15
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Best Local Similarity
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2 TPRPLPLI 9
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US-08-336-343A-29
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US-09-086-168B-3
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Gaps
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APPLICANT: Akerstrom, Claes
APPLICANT: Rask, Lars
APPLICANT: Rask, Lars
APPLICANT: Crumley, Garegg R.
APPLICANT: Morse, Clarence C.
APPLICANT: Morse, Clarence C.
APPLICANT: Murray, Edward M.
APPLICANT: Hjalm, Goran
TITLE OF INVENTION: Thereof and DNA Encoding Same
NUMBER OF SEQUENCES: 106
CORRESPENDENCE ADDRESS:
ADDRESSEE: Rhone-Poulenc Rorer Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         .
0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                    SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/336,343A FILING DATE: 08-NOV-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3: Rhone-Poulenc Rorer Inc. 500 Arcola Rd., 3C43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US95/15203
FILING DATE: 22-NOV-1995
PRIOR PPLICATION DATA:
APPLICATION NUMBER: US 08/344,836
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: Macintosh
OPERATING SYSTEM: System 7.5.1
SUFWARE: Word 6.0 (Patentin)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/652,877
                                                                                                                                                                                                                                                                      NAME: Coruzzi, Laura A.
REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 7683-065
TELECOMUNICATION INFORMATION:
TELEPHONE: (212) 790-9090
TELEFAX: (212) 869-9741/8864
STATE: New York
COUNTRY: U.S.A.
ZIP. 10036-2711
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPALIBLE
COMPUTER: IBM PC COMPALIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 29, Application US/08652877
Patent No. 6187548
                                                                                                                                                                                                                                                                                                                                                           TELEX: (212) 00.
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 29:
SEQUENCE CHARACTER.STICS:
LENGTH: 11 amino acids
TYPE: amino acids
TYPE: amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ZIP: 19426-0107
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                 FILING DATE: 08-NOV-1994
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 100..
Society of Stative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-08-336-343A-29
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 7
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TYPE: PRT; ORGANISM: Caenorhabditis elegans: US-09-845-917A-17
                                                                                     TELEFAX: 610-454-3808
INFORMATION FOR SEQ ID NO: 29:
SEQUENCE CHARACTERISTICS:
LENGTH: 11 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 100.C
Matches 6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                          : peptide
NO
                                                                                                                                                                                                                                                                   internal
                                                                                                                                                                                                . Jeulogy: linear

; MOLECULE TYPE: peptid

; HYPOTHETICAL: NO

; FRAGMENT TYPE: internation 18-08-476-515A-29
                                                                                                                                                                  TYPE: amino acid STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            9 PRPLPV 14
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LENGTH: 11
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0
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                                                                                                                                                                                                                                                                                                                                                                                                                         43.0%; Score 34; DB 3; Length 11; 100.0%; Pred. No. 36; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: Martin Savitzky
STREET: Rhone-Poulenc Rorer Inc., 500 Arcola Rd.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: WO PCT/SE94/00483 FILING DATE: 24-MAY-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: SE 9301764-8
FILING DATE: 24-MAY-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: 07-JUN-1995
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION DATA:
                                                                                                                                     A1355E-US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US 08/344,836 FILING DATE: 23-NOV-1994 FALOR PRICK APPLICATION DATE: APPLICATION NUMBER: WO PCT/SE94/01
                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/487,314
FILING DATE: 07-JUNE-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 29, Application US/08476515A Patent No. 6239270 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: Compaq PC
OPERATING SYSTEM: Windows 95
SOFTWARE: Word 7.0 (Patentin)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/476
                                                                                           NAME: Savitzky, Martin
REGISTRATION NUMBER: 29,699
REFERENCE/DOCKET NUMBER: A135:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 610-454-3816
TELEPHONE: 610-454-3808
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Crumley, Gregg R. APPLICANT: Worse, Clarence C APPLICANT: Murray, Edward M. APPLICANT: Hjalm, Goran
                                                                                                                                                                                                              INFORMATION FOR SEQ ID NO: 29: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Akerstrom, Goran
APPLICANT: Juhlin, Claes
APPLICANT: Rask, Lare
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Floppy disk
FILING DATE: 23-NOV-1994
                                                                                                                                                                                                                                                     LENGTH: 11 amino acids
TYPE: amino acid
STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 100.
5. 6; Conservative
                                                                                                                                                                                                                                                                                                                                                                    ; FRAGMENT TYPE: internal US-08-652-877-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ZIP: 19426-0107
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy o
                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE: peptide HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
ADDRESSEE: Martin Sa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Collegeville
                                                                                                                                                                                                                                                                                                             linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          9 PRPLPV 14
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| RESERVATION WADER: 29,599
| RESERVATION WADER: 29,599
| RESERVATION WADER: 29,599
| RESERVATION WADER: 20,590
| RESERVATION WADER: 20,5044-306.
| RESERVATION FOR SEQ. ID NO. 29, 10-444-306.
| RESERVATION FOR SEQ. ID NO. 29, 10-444.
| RESE
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us-10-009-049-2.szlm15.rai

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FILING DATE: 16-FEB-1996
ATTORNEY/AGENT INFORMATION:
NAME: MABTORY, S. LESLIE
REGISTRATION NUMBER: 18,872
REFERENCE/DOCKET NUMBER: 1101.
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: 16-FEB-1996
CLASSIFICATION: 435
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LENGTH: 15 amino acids
                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
LENGTH: 12 amino acids
                                                                                                                                                                                                                                                                    TOPOLOGY: unknown

MOLECULE TYPE: peptide
US-09-500-124-255
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               , MOLECULE TYPE: peptide US-08-602-999A-410
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                                                                                                                                                                                                                                                   amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: KAY, Brian K.
APPLICANT: THORN, Judith M.
APPLICANT: QUILLIAM, Lawrence A.
APPLICANT: FOWLKES, Dana M.
APPLICANT: FOWLKES, Dana M.
APPLICANT: RIDER, James E.
TITLE OF INVENTION: SH3 BINDING PEPTIDES AND METHODS OF TITLE OF INVENTION: ISOLATING AND USING SAME
NUMBER OF SEQUENCES: 467
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Indels
                                                                                                       COMPUTER READABLE FORM:

MEDTUM TYEE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.30
SOFTWARE: PATENTIN BATA:
APPLICATION NUMBER: US/08/602,999A
FILING DATE: 16-FEB-1996
FLING DATE: L6-FEB-1996
FLING DATE: L6-FEB-1996
FLING DATE: L6-FEB-1996
FLING TOANION: WABER: US/08/02
FREERRATION NUMBER: 18,872
REFERRATION NUMBER: 1101-202
REFERRATION NUMBER: 130-9090
TELLEPAN: (212) 869-9741/8864
TELLEPAN: (212) 869-9741/8864
TELLEPAN: SEQ ID NO: 255:
SEQUENCE CHARACTERISTICS:
LENGTH: 2 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/500,124
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%; Prec. ...
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSEE: Pennie & Edmonds
STREET: 1155 Avenue of the Americas
CITY: New York
                         1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 255, Application US/09500124
Patent No. 6432920
GENERAL INFORMATION:
APPLICANT: SPARKS, Andrew B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/602,999
        Pennie & Edmonds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 100.
Matches 6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY: unknown MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STATE: New York COUNTRY: U.S.A. ZIP: 10036-2711
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   9 PRPLPV 14
                                                                                    COUNTRY: U.S.A. ZIP: 10036-2711
                                                                    New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3 PRPLPV 8
                                               New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-08-602-999A-255
        ADDRESSEE:
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Gaps
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                                                                                                                                                                                                                                                                                                                                              Query Match 43.0%; Score 34; DB 4; Length 12; Best Local Similarity 100.0%; Pred. No. 39; Matches 6; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: SPARKS, Andrew B.
APPLICANT: KAY, Brian K.
APPLICANT: HORN, Judith M.
APPLICANT: OUILLIAM, Lawrence A.
APPLICANT: DER, Channing J.
APPLICANT: POWLKES, Dana M.
APPLICANT: FUBER, James B.
TITLE OF INVENTION: SE3 BINDING PEPTIDES AND METHODS OF
TITLE OF INVENTION: 1813 BINDING AND USING SAME
NUMBER OF SEQUENCES: 467
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/602,999A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
ADDRESSEE: Pennie & Edmonds
STRERT: 1155 Avenue of the Americas
CITY: New York
COUNTRY: U.S.A.
ZIP: 10036-2711
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
COMPUTER: IBM PC Compatible
COMPUTER: IBM PC Compatible
COMPUTER: IBM PC COMPATIBLE
COMPUTER: PC PLOSS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1101-202
1101-202
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 410, Application US/08602999A Patent No. 6184205 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY AGENT INFORMATION:
NAME: Misrock, S. Leslie
REGISTRATION NUMBER: 18,872
REFERENCE/DOCKET NUMBER: 1101-
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 790-9090
TELEFAX: (212) 869-9741/8864
                                             TELEPHONE: (212) 790-9090
TELEFAX: (212) 869-9741/8864
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 255:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 410:
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APPLICANT: KAY, Brian K.
APPLICANT: THORN, Judith M.
APPLICANT: QUILLIAM, Lawrence A.
APPLICANT: DER, Channing J.
APPLICANT: FOWLES, Dana M.
APPLICANT: RIDER, James E.
TITLE OF INVENTION: SH3 BINDING PEPTIDES AND METHODS OF
TITLE OF INVENTION: ISOLATING AND USING SAME
NUMBER OF SEQUENCES: 467
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            43.0%; Score 34; DB 4; Length 15; 100.0%; Pred. No. 49;
43.0%; Score 34; DB 3; Length 15; 100.0%; Pred. No. 49; 1. Indels 1. Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/500,124
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1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1101-202
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION:
PRIOR APPLICATION:
APPLICATION NUMBER: 08/602,999
FILING DATE: 16-FEB-1996
ATTORNEY/AGENT INFORMATION:
NAME: Misrock, S. Leslie
REGISTRATION NUMBER: 1101-202
REFERENCE/DOCKET NUMBER: 1101-202
TELECOMMUNICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 14
US-08-602-999A-86
; Sequence 86, Application US/08602999A
; Patent No. 6184205
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                   Sequence 410, Application US/09500124 Patent No. 6432920
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (212) 869-9741/8864
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEX: 66141 PENNIE INFORMATION FOR SEQ ID NO: 410: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (212) 790-9090
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity 100.
Matches 6; Conservative
    Query Match
Best Local Similarity 100.
Matches 6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ZIP: 10036-2711
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 amino acid
                                                                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: SPARKS,
APPLICANT: KAY, Bri
APPLICANT: THORN, O
APPLICANT: QUILLIAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         9 PRPLPV 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STATE: New York COUNTRY: U.S.A.
                                                                                 9 PRPLPV 14
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                                                                                                                   4 PRPLPV 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New York
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-09-500-124-410
                                                                                                                                                                              RESULT 13
US-09-500-124-410
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX:
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Gaps
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APPLICANT: SPARKS, ANDREW B.
APPLICANT: THORN, JUDITH M.
APPLICANT: QUILLIAM, LAWRENCE A.
APPLICANT: QUILLIAM, LAWRENCE A.
APPLICANT: DER, CHANNING J.
TITLE OF INVENTION: Src SH3 BINDING PEPTIDES AND METHODS OF
TITLE OF INVENTION: ISOLATING AND USING SAME
NUMBER OF SEQUENCES: 106
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: OBLON, SPIVAK, MCCLELLAND, MAIER & NEUSTADT, ADDRESSEE: P.C. STREET: 1755 S. Jefferson Davis Highway, Suite 400 CITY: Arlington
                                           TITLE OF INVENTION: SAME E.

TITLE OF INVENTION: 1SOLATING PEPTIDES AND METHODS OF TITLE OF INVENTION: 1SOLATING AND USING SAME NUMBER OF SEQUENCES: 467

CORRESPONDENCE ADDRESS: ADDRESSEE: Penning STRPPM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Indels
                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/602,999A FILING DATE: 16-FEB-1996
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Misrock, S. Leslie
REGISTRATION NUMBER: 18,872
REFERENCE/DOCKET NUMBER: 1101-202
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    41.8%; Score 33; DB 3; 62.5%; Pred. No. 59;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             3; Mismatches
                                                                                                                                                                                          STREET: 1155 Avenue of the Americas CITY: New York
                                                                                                                                                                                                                                                                           COUNTRY: U.S.A.
ZIP: 10036-2711
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMpatible
COMPUTER: FROM PC-DOS/MS-DOS
SPARKS, Andrew B.
KAY, Brian K.
THORN, Judith M.
QUILLIAM, Lawrence A.
DER, Channing J.
FOWLKES, Dana M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 86, Application US/08278865
Patent No. 6303574
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: (212) bv. TELEFAX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 86:
SEQUENCE CHARACTERISTICS:
TENGTH: 13 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 62.5
Matches 5; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: unknown
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                   New York
: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STATE: Virginia COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-08-602-999A-86
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-08-278-865-86
                                       APPLICANT
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Query Match 41.8%; Score 33; DB 4; Length 13; Best Local Similarity 62.5%; Pred. No. 59; Matches 5; Conservative 3; Mismatches 0; Indels
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: U$/08/278,865
TLING DATE:
CLASSIFICATION: 514
ATTONEY/AGENT INFORMATION:
NAME: Villacorta, Gilberto M.
REGISTRATION NUMBER: 34,038
REFERENCE/DOCKET NUMBER: 4980-007-0
TELEPHONE: (703) 413-320
TELEPHONE: (703) 413-320
TELEFAX: 248855 OPAT UR
SECUROMOTICATION INFORMATION:
TELEFAX: 248855 OPAT UR
SECUROMOTICATION OF 86:
SEQUENCE CHARACTERISTICS:
LENGTH: 3 amino acids
TYPE: amino acid
TYPE: amino acid
US-08-278-865-86
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Search completed: April 2, 2004, 10:12:45 Job time : 15 Becs

8 SPRPLPVV 15 :|||||:: 2 TPRPLPML 9

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; 0

Gaps 0;

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GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
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- protein search, using sw model OM protein April 2, 2004, 10:05:03; Search time 11 Seconds (without alignments) 131.170 Million cell updates/sec Run on:

US-10-009-049-3 82 1 CRAHSFVSPRPLPVV 15

Title: Perfect score: Sequence:

BLOSUM62 Gapop 10.0 , Gapext 0.5 Scoring table:

283366 segs, 96191526 residues Searched:

Total number of hits satisfying chosen parameters:

2522

Minimum DB seq length: 0 Maximum DB seq length: 15

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database:

PIR 78:*
1: pir1:*
2: pir2:*
3: pir3:*
4: pir4:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

		hypothetical prote	photosystem II 3.7	neurotoxin-associa	orf 1 rara 5'-regi	sin	protein QF200037 -	Ig kappa chain V-I	aggrecan - bovine		seed protein ws-18	photosystem II pro	ic	hypothetical prote	inas	gentisate 1,2-diox	mixed lymphocyte r	triacylglycerol li	protein QF200053 -	deoxynucleotidyltr	orf19 3' of eryK -	porphobilinogen sy	xylose isomerase (angiotensin-conver	FMRFamide-like pro	bradykinin-potenti	substance P - chic	ı	substance P - Atla	probable substance
	ID	S27140		I44644	S15594	XAVI9B	PA0060		S42620	XGHUE	C61497	801119	JH0460	836668			PX0031	: PX0030			·	S62641	2 A31576	XAVIGB	2 A43977	XASNBA	2 JN0023	82	S23	\$33300
	Length DB	14 2	7								7		3	13 2	4		Ŋ	0									11			11
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	Score		25	24	23	23	23	22	21	21	21	21	21	21	21	21	21		20		20	20	19	19	19	19	19	19	19	19
1000	No.		7	m	4	5	9	7	œ	6		11		13		15		17	18	19			22				26		28	29

bradykinin-potenti	bradykinin-potenti	protein QF200070 -	Ig H chain V-D-J r	hypothetical 1.5K	dichloromethane de	dichloromethane de	protein QA100006 -	beaded-chain filam	hypothetical prote	ribosomal protein	enamelin f - bovin	Ig H chain V-D-J r	sphingomyelinase -	matrix metalloprot	alcohol dehydrogen
C37196	D37196	PA0104	PH1614	B39109	A60929	B60929	PA0027	A32921	B32800	PA0098	S10783	PH1618	C39745	S71948	866196
7	~	~	7	7	7	~	7	7	7	~	N	~	7	~	7
11	11	14	14	15	15	15	15	15	15	12	c o	c 0	10	10	11
23.2	23.2	23.2	23.2	23.2	23.2	23.2	23.2	23.2	23.2	22.6	22.0	22.0	22.0	22.0	22.0
19	19	19	19	19	19	19	19	19	19	18.5	18	18	18	18	18
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ALIGNMENTS

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Cispecies: Homo sapiens (man)
Cibate: 05-Mar-1994 #sequence_revision 12-May-1995 #text_change 18-Aug-2000
Cibate: 05-Mar-1994 #sequence_revision 12-May-1995 #text_change 18-Aug-2000
Ciracession: S27140
Ciracession: S27140
DNA Seq. 2, 347-358, 1992
A;Title: Sequence analysis of the 5' flanking region of the human estrogen receptor gene. A;Reference number: S27140; MUID: 93075998; PMID: 1476547
A;Reference number: S27140
A;R
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ô
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 35.4%; Score 29; DB 2; Length 14; 54.5%; Pred. No. 1.40+02; tive 1; Mismatches 4; Indels
hypothetical protein 1 estrogen receptor 5'-region - human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity 54.5
Matches 6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2 RAHSFVSPRPL 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     임
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RESULT 2

photosystem II 3.7% protein - spinach (fragment)
Cispecies: Spinacia oleracea (spinach)
Cispecies: Spinacia oleracea (spinach)
Cibate: 01-Dec-1989 #sequence_revision 01-Dec-1989 #text_change 18-Jun-1993
CiAccession: S01122
Rischroeder, W.P.; Henrysson, T.; Akerlund, H.E.
FEBS Lett. 235, 289-292, 1988
A;Title: Characterization of low molecular mass proteins of photosystem II by N-terminal A;Reference number: S01120
A;Accession: S01122

A,Molecule type: protein A,Residues: 1-12 <SCH> C,Keywords: chloroplast; photosynthesis; photosystem II; thylakoid

Gaps . 0 Query Match
30.5%; Score 25; DB 2; Length 12;
Best Local Similarity 57.1%; Pred. No. 5.5e+02;
Matches 4; Conservative 3; Mismatches 0; Indels

.; 0

5 AYAFLSP 11 3 AHSFVSP 9 à qq

RESULT 3 144644

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Matches
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S42620
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A, Residues: 1-13 <OND>
A, Note: the structure of the peptide was confirmed by synthesis
A, Onement: This peptide also potentiates bradykinin by inhibiting the kinases that inact
C, Superfamily: bradykinin-potentiating peptide
C, Superfamily: bradykinin-potentiating peptide
C, Keywords: angiotensin-converting enzyme inhibitor; pyroglutamic acid
F;1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              C. Accession: S15594

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J. Protein Chem. 10, 415-425, 1991
A;Title: Clostridium botulinum types A, B, C1, and E produce proteins with or without A;Reference number: A44644; MUID:92143938; PMID:1781887
A;Contents: type B
neurotoxin-associated protein type B Hn+ 57K chain - Clostridium botulinum (fragment) C;Species: Clostridium botulinum C;Date: 07-Oct-1994 #sequence_revision 07-Oct-1994 #text_change 07-Oct-1994 C;Accession: I44644
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C;Date: 13-Jul-1981 #sequence_revision 13-Jul-1981 #text_change 08-Dec-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C;Species: Homo sapiens (man)
C;Date: 04-Jun-1999 #sequence_revision 04-Jun-1999 #text_change 28-Jun-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  28.0%; Score 23; DB 4; Length 9; 60.0%; Pred. No. 2.8e+05; tive 2; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Molecule type: protein
A;Residues: 1-10 <SOM>
A;Note: sequence extracted from NCBI backbone (NCBIP:83783)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  angiotensin-converting enzyme inhibitor V-9 - jararaca
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          C; Keywords: hemagglutinin
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Best Local Similarity
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Matches 5; Conserv
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A; Status: preliminary
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C;Species: Fusarium sporotrichioides
C;Date: 20 Feb-1995 #sequence_revision 20-Feb-1995 #text_change 23-Mar-2001
C;Accession: PA0060
R;Chow, L.P.; Fukaya, N.; Sugiura, Y.; Ueno, Y.; Tabuchi, K.; Tsugita, A.
submitted to JIPID. October 1994
A;Description: Two dimensional polyacrylamide gel electropheresis of Fusarium sporotrich:
A;Reference number: PA0051
A;Reference number: PA0050
A;Reference number: PA0050
A;Residues: 1-15 CCHO>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ig kappa chain V-II region (anti-myelin-associated glycoprotein, PEC) - human (fragment) C;Species: Homo sapiens (man) C;Species: Homo sapiens (man) C;Species: Homo sapiens (man) C;Species: Homo sapiens (man) C;Species: Homo E.0157; C61458 C;Accession: PLO157; C61458 C;Accession: PLO157; C61458 C;Accession: PLO157; C61458 C;Accession: R;Brouet, J.C.; Dellagi, K.; Gendron, M.C.; Chevalier, A.; Schmitt, C.; Mihaesco, E. A;Title: Expression of a public idiotype by human monoclonal IgM directed to myelin-assoca; A;Reference number: A61458; MUID:90039128; PMID:2478651
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A;Residues: 1-13 <BR2>
C;Comment: This protein is one of monoclonal IgM reactive with myelin-associated glycopro
C;Keywords: glycoprotein; heterotetramer; immunoglobulin
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C;Species: Bos primigenius taurus (cattle)
C;Date: 19-Mar.1997 #sequence_revision 24-Jul-1998 #text_change 24-Jul-1998
C;Accession: S42620
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Score 23; DB 1; Length Lo. Pred. No. 1.3e+03;
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A;Title: Aggrecan in bovine tendon.
A;Feference number: $42620; MUID:94340214; PMID:7520336
A;Accession: $42620
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Pred. No. 1.5e+03;
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100.0%; Pred. No. ...
                                                                             1; Mismatches
                        28.0%;
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Best Local Similarity 100...
                                                                             4; Conservative
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                                                                                                                                 9 PRPLPVV 15
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                           Query Match
Best Local Similarity
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Best Local Similarity
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A, Molecule type: protein
A, Residues: 1-13 - HUJ>
A, Experimental source: bone marrow
A, Experimental source: bone marrow
A, Experimental source: bone marrow
C, Comment: This peptide belongs to a family of Cys-rich, carionic peptides of low molecul
C, Comment: This peptide has antimicrobial activity by a non-oxygen-dependent mechanism.
C, Superfamily: unassigned animal peptides
F; S, Disulfide bonds: interchain (to 13) #status experimental
F; T, Disulfide bonds: interchain (to 7) #status experimental
F; 11, Disulfide bonds: interchain (to 5) #status experimental
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       R.Hu. J.; Bennett, H.P.J.; Lazure, C.; Solomon, S.
Biochem. Biophys. Res. Commun. 180, 558-565, 1991
A,Tille: Isolation and characterization of corticostatic peptides from guinea pig bone ma
A,Reference number: JH0458; MUID:92062075; PMID:1659400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   hypothetical protein 133 - Plectonema boryanum (fragment)
C;Species: Plectonema boryanum
C;Species: Plectonema boryanum
C;Dacession: S3668
R;Fujita, Y.; Matsumoto, H.; Takahashi, Y.; Matsubara, H.
submitted to the RMBL Data Library, August 1992
A;Description: Identification of the nifDK-like gene (ORF467) involved in the biosynthesi
A;Reference number: S36668
                      FEBS Lett. 235, 283-288, 1988
A,Title: Identification of a new gene in the chloroplast genome encoding a low-molecular-
A,Reference number: S01119
                                                                                                                                                                                                                                                                   C;Superfamily: photosystem II protein psbK
C;Keywords: chloroplast; membrane-associated complex; photosynthesis; photosystem II; thy
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C;Species: Cavia porcellus (guinea pig)
C;Date: 30-Jun-1992 #sequence_revision 30-Jun-1992 #text_change 18-Aug-2000
C;Accession: JH0460
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R;Murata, N.; Miyao, M.; Hayashida, N.; Hidaka, T.; Sugiura, M.
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                                                                                                                                                                                                                                                                                                                   25.6%; Score 21; DB 2; I
75.0%; Pred. No. 2.7e+03;
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Best Local Similarity 42.9
Matches 3; Conservative
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Best Local Similarity
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Best Local Similarity
Matches 3: Conserv
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A;Molecule type: DNA
A;Residues: 1-13 <FUJ>
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A.Title: Microsequence analygis of winged bean seed proteins electroblotted from two-dim A,Fitle: mumber: A61491; MUID:89351606; PMID:2765119
A,Accession: C61497
A,Status: preliminary
A,Molecule type: protein
A,Residues: 1-11 cHIR>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A,Molecule type: protein
A,Residues: 1-10 <WEI>
C;Comment: The identity of the glycoprotein from which this peptide is derived is unknow
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              R;Weiss, J.B.; Lote, C.J.; Bobinski, H. Nature New Biol. 234, 25-26, 1971
A;Title: New How molecular weight glycopeptide containing triglucosylcysteine in human A;Reference number: A03187; MUID:72034940; PMID:5286858
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0
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C;Species: chloroplast Spinacia oleracea (spinach)
C;Date: 30-Jun-1989 #sequence_revision 31-Dec-1990 #text_change 19-Jan-1996
C;Accession: S01119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              C;Species: Psophocarpus četragonolobuš (winged bean)
C;Datts: 07-Oct-1994 #sequence_revision 07-Oct-1994 #text_change 07-Oct-1994
C;Accession: C61497
R;Hirano, H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           C;Species: Homo sapiens (man)
C;Date: 20-Jun-2000 #sequence_revision 20-Jun-2000 #text_change 20-Jun-2000
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C;Superfamily: unassigned animal peptides
C;Superfamily: unassigned animal peptides
C;Keywords: glycoprotein
F;1/Binding site: carbohydrate (Cys) (covalent) #status experimental
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     .
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                                                                                                                                   Length 7;
                                                                                                                                Score 21; DB 2; I Pred. No. 2.8e+05;
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                                                                                                                                                                                                                                                                                                                                                                                                                               erythrocyte membrane glycopeptide - human
  A,Molecule type: protein
A,Residues: 1-7 <VOG>
A;Experimental source: flexor tendon
C;Keywords: cartilage
                                                                                                                                   25.6%;
57.1%;
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                                                                                                                                   Query Match
Best Local Similarity 57.1
Matches 4; Conservative
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Matches 3; Conserva
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CRAHSFVSPRP 11

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gentisate 1,2-dioxygenase (EC 1.13.11.4) - Comamonas testosteroni (fragment)
C;Species: Comamonas testosteroni
C;Date: 10-Aug-1990 #sequence_revision 10-Aug-1990 #text_change 23-Jun-1993
C;Accession: A35232
R;Harpel, M.R.; Lipscomb, J.D.
B;Ol. Chem. 265, 6310-6311, 1990
A;Title: Gentisate 1,2-dioxygenase from Pseudomonas. Purification, characterization, and A;Reference number: A35232, MUID:90202907; PMID:2156846
                                                                                                                         protein kinase (EC 2.7.1.37) - starfish (Pisaster ochraceus)
C;Species: Pisaster ochraceus
C;Species: Pisaster ochraceus
C;Date: 19-Mar-1997 #sequence_revision 19-Mar-1997 #text_change 01-Aug-1997
C;Accession: 512904
R;Sanghera, J.S.; Aebersold, R.; Morrison, H.D.; Bures, E.J.; Pelech, S.L.
RESS Lett. 273, 223-226, 1990
A;Title: Identification of the sites in myelin basic protein that are phosphorylated by A;Reference number: 512904; MUID:91032186; PMID:1699809
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25.6%; Score 21; DB 2; Length 15;
Best Local Similarity 50.0%; Pred. No. 3.1e+03;
Matches 4; Conservative 1; Mismatches 3; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                       A Status: preliminary
A,Molecule type: protein
Residues: 1-14 < SAN>
C;Keywords: phosphotransferase
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Best Local Similarity 44.41
Matches 4; Conservative
2 CEGRNRTIPSP 12
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A;Molecule type: protein
A;Residues: 1-15 <HAR>
C;Keywords: oxidoreductase
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GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
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OM protein - protein search, using sw model

April 2, 2004, 09:58:38; Search time 7.66667 Seconds (without alignments) 101.876 Million cell updates/sec Run on:

US-10-009-049-3 82 1 CRAHSFVSPRPLPVV 15 Title: Perfect score:

BLOSUM62 Gapop 10.0 , Gapext 0.5 Scoring table: Sequence:

141681 segs, 52070155 residues Searched:

801 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 15

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

SwissProt_42:* Database :

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

STIMMARTES

		Description	P80635 zea mays (m	P43172 ascaris suu		_	_		periplane	P02728 homo sapien							manduca s	bothrops	Ω	agkist	P19850 gallus gall	gadus				_		~	9	P80940 lymantria d	20 rattus	8497]	0622 zea may	40751 u	P19425 aedes aegyp
SUMMARIES	;	OI.	UC29 MAIZE	FAR9 ASCSU	MK1 PALPR		PPK4_PERFU	BPP1_BOTJA	PPK4_PERAM	GLEM HUMAN		TIN2 HOPTI	TIN3_HOPTI	MGMT_BOVIN	XYLA_STRSQ	BPP2 BOTJA	FARP MANSE	BPP3_BOTIN	BPP4_BOTIN	BPPB_AGKHA	TKNA CHICK	TKNA GADMO	TKNA_ONCMY	TKNA_SCYCA	ECDE_LYMDI	UC15_MAIZE	CX3B_CONQU	TEMD_RANTE	UPS1_UPEIN	ECDC_LYMDI	C1QA_RAT	LMA2_LOCMI	UC16_MAIZE	TKU1_UREUN	TMOF_AEDAE
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	RESULT PROPERTY OF SECUL STATE OF SECUL SE	SUL

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PPK4 PERFU
P82690;
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Chernysh S., Cociancich S., Briand J.-P., Hetru C., Bulet P.;
"The inducible antibacterial peptides of the hemipteran insect
Palomena prasina: identification of a unique family of proline-rich
peptides and of a novel insect defensin.";
J. Insect Physiol. 42:81-89(1996).
-!- FUNCTION: Antibacterial peptide active against Gram-negative
        Peptides 16:491-500(1995).
-!- SIMILARITY: Belongs to the FARP (FWRFamide related peptide)
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                                                                                                                                                                                                                                                                           Palomena prasina (Green shield bug).
Bukaryota, Metazoa, Arthropoda, Hexapoda, Insecta, Pterygota,
Neoptera, Paraneoptera, Hemiptera, Buhemiptera, Heteroptera,
Panheteroptera, Pentatomomorpha, Pentatomoidea, Pentatomidae,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Palomena prasina (Green shield bug).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Neoptera; Paraneoptera; Hemiptera; Euhemiptera; Heteroptera;
Panheteroptera; Pentatomomorpha; Pentatomoidea; Pentatomidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      29.3%; Score 24; DB 1; Length 15; 80.0%; Pred. No. 5.3e+02; ive 0; Mismatches 1; Indels
                                                                                                          1; Indels
                                                                                     Score 24; DB 1; Length 9; Pred. No. 1.4e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Antibiotic; Insect immunity. SEQUENCE 15 AA; 1838 MW; 21407E663CE46299 CRC64;
                                                    AMIDATION.
524F073774176877 CRC64;
                                                                                                                                                                                                                                01-NOV-1995 (Rel. 32, Created)
01-NOV-1995 (Rel. 32, Last sequence update)
28-FEB-2003 (Rel. 41, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-NOV-1995 (Rel. 32, Created)
01-NOV-1995 (Rel. 32, Last sequence update)
28-FEB-2003 (Rel. 41, Last annotation update)
Metalnikowin IIA.
                                                                                                                                                                                                               15 AA.
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66.7%;
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9 AA; 1012 MW;
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                                       Neuropeptide; Amidation.
MOD_RES 9 9
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                                                                                                                                7 VSPRPL 12
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                                                                                                                                                      2 LGPRPL 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NCBI_TaxID=55431;
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28-FEB-2003 (Rel
Metalnikowin I.
 Ascaris suum.";
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P80409;
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P80408;
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the American cockroach.";
J. Comp. Neurol. 419:352-363(2000).
-!- FUNCTION: Mediates visceral muscle contractile activity (myotropic
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Palomena prasina: identification of a unique family of proline-rich peptides and of a novel insect defensin.";
J. Insect Physiol. 42.81-89(1996).
-!- FUNCTION: Antibacterial peptide active against Gram-negative
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21-UUJ-1996 (Rel. 01, Created)
1-FBE-1994 (Rel. 28, Last sequence update)
28-FBB-2003 (Rel. 41, Last annotation update)
Bradykinin-potentiating peptide S3,1 (13A) (Angiotensin-converting
enzyme inhibitor V-9)
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Bothrops insularis (Island jararaca) (Queimada jararaca).
Bukaryota; Metazoa, Chordata; Craniata; Vertebrata; Euteleostomi;
Lepidosauria; Squamata; Scleroglossa; Serpentes; Colubroidea;
Viperidae; Crotalinae; Bothrops.
NCBI_TaxID=8724, 8723;
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Neoptera, Orthopteroidea, Dictyoptera, Blattaria, Blattoidea,
                                                                                                                                                                                                                                                                  29.3%; Score 24; DB 1; Length 15; 80.0%; Pred. No. 5.3e+02; 1ive 0; Mismatches 1; Indels
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                                                                                                                                                                          Antibiotic; Insect immunity. SEQUENCE 15 AA; 1893 MW; 23835D063B946299 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  activity).
-!- MASS SPECTROMETRY: MW=1437.9; METHOD=MALDI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   16-OCT-2001 (Rel. 40, Created)
16-OCT-2001 (Rel. 40, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
Pyrokinin-4 (PEF-PK-4) (YXPRL-amide).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Periplaneta fuliginosa (Smokybrown cockroach)
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                                                                                                                    bacteria.
-!- INDUCTION: By bacterial infection.
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MEDLINE=20189894; PubMed=10723010;
Predel R., Eckert M.;
                                                                                                                                                                                                                                                                                                                                 4; Conservative
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NCBI_TaxID=36977;
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Pred. No. 9e+02;
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16-OCT-2001 (Rel. 40, Last sequence update)
10-OCT-2003 (Rel. 42, Last annotation update)
                                                                                                                                                                                                           21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
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                                                                                                                                                                          10 AA
                   0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                tigerina.";
T Riol. Chem. 276:2701-2707(2001).
                                                                                                                                                                            PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRT;
                                                                                                                                                                                                                                                               Erythrocyte membrane glycopeptide.
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 57.1%;
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Glycoprotein; Erythrocyte.
CARBOHYD 1 1
                   4; Conservative
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                                                     4 HSFVSPR 10
                                                                                      5 HDVYSPR 11
     Best Local Similarity
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Best Local Similarity
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GLEM HUMAN
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TIN4 HOPTI
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                                                                                                                                                                                   SPECIES-B.insularis; TISSUE=Venom;
MEDLINE=90351557; PubMed=2386615;
Cintra A.C.O., Valeira C.A., Giglio J.R.;
Primary Structure and biological activity of bradykinin potentiating peptides from Bothrops insularis snake venom.";
J. Protein Chem. 9:221-227(1990)
-!-FUNCTION: This peptide both inhibits the activity of the angiotensin-converting enzyme and enhances the action of bradykinin by inhibiting the kinases that inactivate it.
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                                                                                                    "Angiotensin-converting enzyme inhibitors from the venom of Bothrops
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDLINE=20189894; PubMed=10723010;
Predel R., Eckert M.;
"Tagma-specific distribution of FXPRLamides in the nervous system of
                              SPECIES=B.jararaca; TISSUB=Venom;
MEDLINE=72118526; PubMed=4334402;
Ondetti M.A., Williams N.J., Sabo E.F., Pluscec J., Weaver E.R.,
                                                                                                                        jararaca. Isolation, elucidation of structure, and synthesis.";
Biochemistry 10:4033-4039(1971).
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Periplaneta americana (American cockroach).
Bukaryota, Metazoa, Arthropoda, Hexapoda, Insecta, Pterygota,
Neoptera, Orthopteroidea, Dictyoptera, Blattaria, Blattoidea,
Blattidae, Periplaneta.
                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 23; DB 1; Length 15; Pred, No. 6.78+02;
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                                                                                                                                                                                                                                                                                                                                                                              Hypotensive agent; Pyrrolidone carboxylic acid.
MOD_RES 1 1 PYRROLIDONE CARBOXYLIC ACID.
SEQUENCE 13 AA, 1388 MW; 6824FC97D83D6774 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     12 12 AMIDATION.
12 AA; 1449 MW; FA7A3049FF42CAAI CRC64;
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-!- SIMILARITY: Belongs to the pyrokinin family.
Neuropeptide; Amidation; Pyrokinin.
MOD EBS 12 AMIDATION.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      16-00T-2001 (Rel. 40, Created)
16-00T-2001 (Rel. 40, Last sequence update)
16-00T-2001 (Rel. 40, Last annotation update)
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IISSUE=Retrocerebral complex;
MEDLINE=99212469; PubMed=10196736;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      28.0%;
57.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      4; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            the American cockroach."
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Best Local Similarity
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Weiss J.B., Lote C.J., Bobinski H.;
"New low molecular weight glycopeptide containing triglucosylcysteine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    in human erythrocyte membrane.";
Nature New Biol. 234:25-26(1971).
-!- PTM: S-linked glycan consists of GIC-GIC-GIC trisaccharide.
-!- MISCELLANEOUS: The identity of the glycoprotein from which this peptide is derived is unknown. No physiological function has been
       Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Buteleostomi,
Mammalia, Butheria, Primates, Catarrhini, Hominidae, Homo.
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Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Amphibia, Batrachia, Anura, Neobatrachia, Ranoidea, Ranidae,
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-!- SUBCELLULULAR LOCATION: Secreted.
-!- TISSUB SPECIFICITY: Skin.
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Pred. No. 1.1e+03;
0; Mismatches 2; Indels
   Indels
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239BFEEAA1F5B1E8 CRC64;
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Devi A.S., Nagaraj R., Sitaram N.;
"Tigerinins: novel antimicrobial peptides from the Indian frog Rana
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PubMed=11031261;
Purna Sai K., Jaganadham M.V., Vairamani M., Raju N.P.,
Devi A.S., Nagarai R., Sitaram N.;
"Tigerinins: novel antimicrobial peptides from the Indian frog Rana
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hoplobatrachus tigerinus (Indian bull frog) (Rana tigerina).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Amphibia, Batrachia, Anura, Neobatrachia, Ranoidea, Ranidae,
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Batrachia; Anura; Neobatrachia; Ranoidea; Ranidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                J. Biol. Chem. 276:2701-2707(2001).
-!- FUNCTION: Antibacterial activity against B.subtilis, B.coli,
S.aureus, M.luteus, P.luulida and S.cerevisiae.
-!- SUBCELLULAR LOCATION: Secreted.
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                                                           25.6%; Score 21; DB 1; Length 11; 75.0%; Pred. No. 1.2e+03; tive 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  25.6%; Score 21; DB 1; Length 12; 75.0%; Pred. No. 1.38+03; tive 1; Mismatches 0; Indels
                  1248 MW; 117D8EFD37605DCB CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AMIDATION.
C2627D8EFD37605D CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        -!- TISSUE SPECIFICITY: Skin.
-!- MASS SPECTROMETRY: MW=1368; METHOD=MALDI.
Amphibian defense peptide; Antibiotic; Amidation.
                                                                                                                                                                                                                                                                                                                                                       16-OCT-2001 (Rel. 40, Created)
16-OCT-2001 (Rel. 40, Last sequence update)
10-OCT-2003 (Rel. 42, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              16-0CT-2001 (Rel. 40, Created)
16-0CT-2001 (Rel. 40, Last sequence update)
10-0CT-2003 (Rel. 42, Last annotation update)
                                                                                                                                                                                                                                                                                                                 12 AA.
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                                                                                                                                                                                                                                                                                                                    PRT;
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                                                                                  Local Similarity 75.0 les 3; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity 75.0
nes 3; Conservative
                                                                                                                                                                                                                                                                                                               STANDARD;
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PubMed=11031261;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                IISSUE=Skin secretion;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NCBI_TaxID=103373;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NCBI_TaxID=103373;
3
11 AA;
                                                                                                                                                     11 PLPV 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            11 PLPV 14
                                                                                                                                                                                               7 PLPI 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Tigerinin-3
                                                                                                                                                                                                                                                                                                            TIN2 HOPTI
P82652;
                                                                                                                                                                                                                                                                                                                                                                                                                                 Tigerinin-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      IIN3 HOPTI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               tigerina."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE
                  SEQUENCE
                                                             Query Match
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DISULFID
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Matches
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                                                                                                        Matches
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Nucleic Acids Res. 18:17-21(1990).
-!- FUNCTION: Repair of alkylated guanine in DNA by stoichiometrically
-transferring the alkyl group at the O-6 position to a cysteine
residue in the enzyme. This is a suicide reaction: the enzyme is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-DEC-1992 (Rel. 24, Created)
01-DEC-1992 (Rel. 24, Last sequence update)
01-OCT-1996 (Rel. 34 Last annotation update)
Methylated-DNA--protein-cyteine methyltransferase (EC 2.1.1.63) (6-O-methylguanine-DNA methyltransferase) (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             irreversibly inactivated.
-!- CATALYTIC ACTIVITY: DNA (containing 6-0-methylguanine) +
[protein]-L-cysteine = DNA (without 6-0-methylguanine) + protein
S-methyl-L-cysteine.
-!- SIMILARITY: WITH SEGMENTS OF E.COLI ADA AND OGT METHYLTRANSFERASE
WHICH ENCOMPASS THE ALKYL-ACCEPTOR RESIDUES.
                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                'Active site amino acid sequence of the bovine O6-methylguanine-DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Bos taurus (Bovine).
Wakaryota: Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Butheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea;
Bovidae; Bovinne; Bos.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ALKYL GROUP ACCEPTOR (BY SIMILARITY)
tigerina.";
J. Biol. Chem. 276:2701-2707(2001).
J. Biol. Chem. 276:2701-2707(2001).
J. FUNCTION: Antibacterial activity against B.subtilis, E.coli, S. aureus, M.luteus, P.putida and S.cerevisiae.
J. SUBCELLULAR LOCATION: Secreted.
J. TISSUE SPECIFICITY: Skin.
J. TISSUE SPECIFICITY: Skin.
J. MASS SPECIFICITY: Skin.
J. MASS SPECIFICITY: Amm=1409; METHOD=MALDI.
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                                                                                                                                                                                                                                                                                                                                                                                                    0; Indels
                                                                                                                                                                                                                                                                                         C0717D8EFD37605D CRC64;
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                                                                                                                                                                                                                                                                                                                                         Score 21; DB 1; I
Pred. No. 1.3e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             9 AA.
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NON TER
                                                                                                                                                                                                                                                                                                                                                                                                 1; Mismatches
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PROSITE; PS00374; MGMT; PARTIAL.
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MEDLINE=90174912; PubMed=2308822;
Rydberg B., Hall J., Karran P.;
                                                                                                                                                                                                                                                                                                                                               25.6%;
                                                                                                                                                                                                                                                      MOD_RES 12 12
SEQUENCE 12 AA; 1411 MW;
                                                                                                                                                                                                                                                                                                                                                                                                          3; Conservative
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Best Local Similarity
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7 PLPI 10
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RESULT 15
FARP_MANSE
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InterFor: IPRO1998; Xylose isom.
PROSITE; PS00172; XYLOSE ISOMERASE 1; PARTIAL.
PROSITE; PS00173; XYLOSE ISOMERASE 2; PARTIAL.
ISOMETASE; PS00173; XYLOSE ISOMERASE 2; PARTIAL.
ISOMETASE; Pentose shunt; Xylose metabolism; Metal-binding; Magnesium.
                                                                                                                                                                              Pawar H.S., Kannan K., Srinivasan M.C., Vartak H.G.; "Purification and characterisation of glucose (xylose) isomerase from Chainia sp. (NCL 82-5-1).";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      "Angiotensin-converting enzyme inhibitors from the venom of Bothrops jararaca. Isolation, elucidation of structure, and synthesis."; Biochemistry 10:4033-4039(1971).
-!- FUNCTION: This peptide both inhibits the activity of the angiotensin-converting enzyme and enhances the action of bradykinin by inhibiting the kinases that inactivate it.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             21-JUL-1986 (Rel. 01, Created)
01-FEB-1994 (Rel. 28, Last sequence update)
28-FEB-2003 (Rel. 41, Last annotation update)
Bradykinin-potentiating peptide 10B (Angiotensin-converting enzyme
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Bothrops jararaca (Jararaca).

Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Lepidosauria; Squamata; Scleroglossa; Serpentes; Colubroidea;
Viperidae; Crotalinae; Bothrops.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TISSUE=Venom;
MEDLINE=72118526; PubMed=4334402;
Ondetti M.A., Williams N.J., Sabo E.F., Pluscec J., Weaver E.R.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             .
0
                                                                              Streptomyces sp. (strain NCL 82-5-1).
Bacteria, Actinobacteria, Actinobacteridae, Actinomycetales,
Streptomycineae, Streptomycetaceae, Streptomyces.
                                                                                                                                                                                                                      Biochem. Biophys. Res. Commun. 155:411-417(1988).
-!- PUNCTION: Involved in D-xylose catabolism.
-!- CATALYTIC ACTIVITY: D-xylose = D-xylulose.
-!- COFACTOR: Binds 2 magnesium ions per subunit (Potential).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PIR; A01255; XAVIGB.

"Hypotensive agent; Pyrrolidone carboxylic acid.

MOD RES

SEQÜENCE 10 AA; 1232 MW; 30C53546C774173 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                              23.2%; Score 19; DB 1; Length 9; 75.0%; Pred. No. 1.4e+05; tive 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                          -!- SUBCELLULAR LOCATION: Cytoplasmic.
-!- SIMILARITY: Belongs to the xylose isomerase family.
PIR; A31576; A31576.
                                                                                                                                                                                                                                                                                                                                                                                                                    9 AA; 983 MW; F64BA1EDC5B87DD1 CRC64;
          01-NOV-1990 (Rel. 16, Created)
01-NOV-1990 (Rel. 16, Last sequence update)
01-OCT-2003 (Rel. 42, Last annotation update)
Xylose isomerase (EC 5.3.1.5) (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         10 AA.
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                                                                                                                                                                    MEDLINE=88326335; PubMed=3415697;
                                                                                                                                                                                                                                                                              SUBUNIT: Homotetramer.
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nes 3; Conservative
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                                                                                                                        NCBI_TaxID=1931;
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6 AHTF 9
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P01022;
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SEQUENCE
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                                                                                                                                                                                                                                                                                                                                           FMRFamide-like neuropeptide.
Manduca sexta (Tobacco hawkmoth) (Tobacco hornworm).
Bukaryota, Metazoa, Arthropoda, Hexapoda, Insecta, Pterygota,
Neoptera, Endopterygota, Lepidoptera; Glossata, Ditrysia, Sphingiodea,
Sphingidae, Sphinginae, Manduca.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     hawkmoth, Manduca sexta.";
Peptides 11:849-856(1990).
-!- FUNCTION: Increases the force of neurally evoked contractions in the major power-producing flight muscles, the dorsal longitudinal muscles and so is likely to play a role in sustaining or promoting flight behavior patterns.
-!- SIMILARITY: Belongs to the FARP (FMRFamide related peptide)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDLINE=91045350; PubMed=2235684;
Kingan T.G., Teplow D.B., Phillips J.M., Riehm J.P., Rao K.R.,
Hildebrand J.G., Homberg U., Kammer A.B., Jardine I., Griffin P.R.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   "A new peptide in the FMRFamide family isolated from the CNS of the
                                                      .
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       Length 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Amidation, Neuropeptide, Pyrrolidone carboxylic acid.
MOD_RES 1 1 PYRROLIDONE CARBOXYLIC ACID.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Indels
                                                   0; Indels
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D3C45229D5B1F2D2 CRC64;
23.2%; Score 19; DB 1; Lv 100.0%; Pred. No. 2.3e+03; iive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                         01-NOV-1990 (Rel. 16, Created)
01-FEB-1994 (Rel. 28, Last sequence update)
28-FEB-2003 (Rel. 41, Last annotation update)
                                                                                                                                                                                                                                                                  10 AA.
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                                                 3; Conservative
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  Query Match
Best Local Similarity
Matches 3; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Search completed: April Job time: 8.66667 secs
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SEQUENCE.
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OM protein - protein search, using sw model

Run on:

April 2, 2004, 10:04:38; Search time 31.3333 Seconds (without alignments) 151.046 Million cell updates/sec

US-10-009-049-3 Title: Perfect score:

1 CRAHSFVSPRPLPVV 15 Sequence: BLOSUM62 Gapop 10.0 , Gapext 0.5 Scoring table:

1017041 seqs, 315518202 residues Searched:

3954 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 15

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

sp_fungi:*
sp_human:*
sp_invertebrate:*
sp_mammal:*
sp_mhc:*
sp_organelle:* sp_vertebrate:*
sp_unclassified:* 8p_rvirus:*
sp_bacteriap:* sp_archea:* sp_bacteria:* sp_rodent:*
sp_virus:* sp plant: * sp phage:* SPTREMBL 25:* 10: Database :

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

sp_archeap:*

	Description	Q95771 ctenosaura	Q95952 sauromalus			Q9r5n3 clostridium		08cq39 rattus norv	Q81v78 polyomaviru	088176 mus musculu	Q7x566 thermus the	Q988z9 psophocarpu	Q9qvi3 cavia (quin	Q9ucc2 homo sapien	000604 homo sapien	Q16220 homo sapien	Ogovto mus musculu
SUMMARIES	ПD	095771	995952	095879	095770	Q9R5N3	Q9TWV0	080639	Q8JV78	088176	Q7X566	Q9S8Z9	O9QVI3	Q9UCC2	000604	016220	OJWQ60
	DB	8	œ	œ	8	7	Ŋ	11	12	1	~	10	11	4	4	4	11
	% Query Match Length Di	15	15	15	15	10	σı	σ	10	13	11	11	13	15	15	σ	δ
	% Query Match	30.5	30.5	30.5	30.5	29.3	28.0	28.0	28.0	28.0	25.6	25.6	25.6	25.6	25.6	24.4	24.4
	Score	25	25	25	25	24	23	23	23	23	21	21	21	21	21	20	20
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RESULT 2 Q95952

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12 11 Q9QVK4 13 4 Q14182 13 4 Q9U452 14 8 Q9G3C0 15 5 Q26ACX8 15 10 Q9S8B1 15 10 Q9S8B1 16 10 Q9CK53 17 10 Q9CK53 18 10 Q9CK54 11 2 Q8CKA1 11 2 Q8CKA1 12 2 Q9CKA1 12 2 Q9CKA1 13 10 Q94SC1 13 10 Q94SC1 13 10 Q94SC1 15 2 Q9CK6 15 2 Q9CK6 16 2 Q9CK6 17 Q9CK6 18 2 Q9CK6 19 2 Q9CK6 19 2 Q9CK6 10 Q94SC1 11 10 Q94SC1 11 10 Q94SC1 12 10 Q94SC1 13 10 Q94SC1 14 Q9CK6 15 10 Q94SC1 16 Q9CK6 17 Q9CK6 18 10 Q94SC1 19 10 Q94SC1 19 10 Q94SC1 10 Q94SC1 11 Q94SC1 12 Q9CK6 13 Q9CK6 14 Q9CK6 15 Q9CK6 16 Q9CK6 17 Q9CK6 18 Q9CK6 19 Q9CK6 19 Q9CK6 10 Q9CK6 10 Q9CK6 11 Q9CK6 11 Q9CK6 12 Q9CK6 13 Q9CK6 14 Q9CK6 15 Q9CK6 16 Q9CK6 17 Q9CK6 18 Q9CK6 19 Q9CK6 19 Q9CK6 10 Q9CK6 10 Q9CK6 10 Q9CK6 10 Q9CK6 11 Q9CK6 11 Q9CK6 12 Q9CK6 13 Q9CK6 14 Q9CK6 15 Q9CK6 16 Q9CK6 17 Q9CK6 18 Q9CK6 18 Q9CK6 18 Q9CK6 19 Q9CK6 19 Q9CK6 19 Q9CK6 10 Q9CK6	9 2 PB3157
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Lepidosauria, Squamata, Iguania, Iguanidae, Iguaninae, Ctenosaura.
NCBI_TaxID=51212,
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                                     01-FEB-1997 (TrEMBLrel. 02, Created)
01-FEB-1997 (TrEMBLrel. 02, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
NADH dehydrogenase subunit 4 (Fragment).
                     15 AA.
                                                                                         Ctenosaura similis (Spiny-tailed iguana). Mitochondrion.
                     PRT;
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Best Local Similarity 80.00
Best Local 4; Conservative
                     PRELIMINARY;
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RESULT 1
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Mitochondrion
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SEQUENCE
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Sites J.W. Jr., Davis S.K., Guerra T., Iverson J.B., Snell H.L.;
"Character congruence and phylogenetic signal in molecular and
morphological data sets: a case study in the living Iguanas (Squamata,
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Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Lepidosauria, Squamata, Iguania, Iguanidae, Phrynosomatinae,
                                                                                                                                                                                        Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Lepidosauria, Squamata, Iguania, Iguanidae, Iguaninae, Sauromalus
NCBI_TaxID=51220,
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30.5%; Score 25; DB 8; Length 15;
Best Local Similarity 80.0%; Pred. No. 1.6e+03;
Matches 4; Conservative 0; Mismatches 1; Indels
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80.0%; Pred. No. 1.6e+03;
tive 0; Mismatches 1; Indels
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01-FEB-1997 (TrEMBLrel. 02, Last sequence update)
01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
NADH dehydrogenase subunit 4 (Fragment).
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01-FEB-1997 (TrEMBLrel. 02, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
NADH dehydrogenase subunit 4 (Fragment).
                                                                                                                                                 Sauromalus varius (San Esteban island chuckwalla).
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15 AA.
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                                                                                                                                                                                                                                                                                                                                                                                       Iguānidae).";
Mol. Biol. Evol. 13:1087-1105(1996).
EMBL; U66233; AAB07480.1; -
GO; GO:0005739; C:mitochondrion; IEA.
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EMBL, U66224; AAB07478.1; -
GO; GO:0005739; C:mitochondrion; IEA.
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PRELIMINARY;
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NCBI_TaxID=43610;
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SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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                                                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Lepidosauria; Squamata; Iguania; Iguanidae; Iguaninae; Cyclura.
NCBI_TaxID=51215;
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Bacteria; Firmicutes; Clostridia; Clostridiales; Clostridiaceae;
Clostridium.
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                                             01-FEB-1997 (TrEMBLrel. 02, Created)
01-FEB-1997 (TrEMBLrel. 02, Last sequence update)
01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            01.MAY-2000 (TrEMBLrel. 13, Created)
01.MAY-2000 (TrEMBLrel. 13, Last sequence update)
01.JUN-2003 (TrEMBLrel. 24, Last annotation update)
Neurotoxin type B HN+ 57 kDa subunit (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           10 AA.
                                                                                                                                 NADH dehydrogenase subunit 4 (Fragment).
                                                                                                                                                                                    Cyclura ricordi (Ricord's rock iguana).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MG]. Biol. Evol. 13:1087-1105(1996).
EMBL, U66237; AAB07473.1; -.
GO, GO:0005739; C:mitochondrion, IEA.
  PRT;
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J. Protein Chem. 10:415-425(1991).
PIR; I44644; I44644.
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Bust Local Similarity 80.v.,
Best Local 4; Conservative
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PRELIMINARY;
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Kawahigashi H., Harada Y., Asano A., Nakamura M.,
"A cis-acting regulatory element that affects the alternative splicing
of a muscle-specific exon in the mouse NCAM gene.";
Biochim. Biophys. Acta 1397:305-315(1998).
EMBL, AB001873; BAA31275.1;
MGD; MGI:97281; Ncaml.
   Viruses; dsDNA viruses, no RNA stage; Polyomaviridae; Polyomavirus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Soiurognathi; Muridae; Murinae; Mus.
                                                                         MEDLINE=21465052; PubMed=11581397;
Bofill-Mas S., Formiga-Cruz M., Clemente-Casares P., Calafell F.,
                                                                                                                                     gastrointestinal tract after exposure to virions or viral DNA."; J. Virol. 75:10290-10299(2001).
BRBL; AF303946; AAM97796.1; -. NON TER 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Bacteria; Deinococcus-Thermus; Deinococci; Thermales; Thermaceae;
                                                                                                       Girones R.;
"Potential transmission of human polyomaviruses through the
                                                                                                                                                                                                                          28.0%; Score 23; DB 12; Length 10; 50.0%; Pred. No. 2.30+03; ative 1; Mismatches 3; Indels
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Pred. No. 3e+03;
1; Mismatches 0; Indels
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NON TER 13 13
SEQÜENCE 13 AA; 1496 MW; CC6098E54C72D732 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                         01-NOV-1998 (TrEMBLrel. 08, Created) 01-NOV-1998 (TrEMBLrel. 08, Last sequence update) 01-OCT-2002 (TrEMBLrel. 22, Last annotation update) Neural cell adhesion molecule (Fragment).
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01-0CT-2003 (TrEMBLrel. 25, Last sequence update)
01-0CT-2003 (TrEMBLrel. 25, Last annotation update)
Hypothetical protein (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                  13 AA.
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MEDLINE=98250618; PubMed=9582442;
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80.0%;
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Query Match
Best Local Similarity 50.0%,
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Best Local Similarity
Matches 4; Conserv
                                               SEQUENCE FROM N.A.
                    NCBI TaxID=10632;
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                                                              STRAIN=USA1;
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                                                                                                                                                                                            Carstensen K., Rinehart K.L., McFarlane I.D., Grimmelikhuijzen C.J., "Isolation of Leu-Pro-Pro-Gly-Pro-Leu-Pro-Arg-Pro-NH2 (Antho-RPamide), an N-terminally protected, biologically active neuropeptide from sea
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Rattus norvegicus (Rat).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Rodentia, Sciurognathi, Muridae, Murinae, Rattus.
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                                                                                                    Eukaryota; Metazoa; Cnidaria; Anthozoa; Zoantharia; Actiniaria; Nynantheae; Actiniidae; Anthopleura.
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MEDLINE=96342418; PubMed=8750786;
Takemura M., Yamauchi K., Yamatodani A.;
"Structural analysis of histamine N-methyltransferase gene.";
Methods Find. Exp. Clin. Pharmacol. 17:1-4(1995).
EMBL; S82579; AAN86745.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   28.0%; Score 23; DB 11; Length 9; 66.7%; Pred. No. 1e+06; tive 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                  28.0%; Score 23; DB 5; Length 9; 80.0%; Pred. No. 1e+06; Live 0; Mismatches 1; Indels
                          01-MAR-2003 (TrEMBLrel. 23, Created)
01-MAR-2003 (TrEMBLrel. 23, Last sequence update)
01-UUN-2003 (TrEMBLrel. 24, Last annotation update)
Histamine N-tele-methyltransferase (Fragment)
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01-0CT-2002 (TrEMBLrel. 22, Last sequence update)
10-0CT-2002 (TrEMBLrel. 22, Last annotation update)
Large T antigen (Fragment).
                                                                                                                                                                                                                                                   Peptides 13:851-857(1992).
SEQUENCE 9 AA; 943 MW; 2908176737686777 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GO; GO:0008168; F:methyltransferase activity; IEA.
GO; GO:0016740; F:transferase activity; IEA.
Transferase; Methyltransferase.
   9 A.A.
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                                                                                      Anthopleura elegantissima (Sea anemone).
   PRT;
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nes 4; Conservative
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Score 21; DB 11; Length 13; Pred. No. 6.7e+03;

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13 AA; 1706 MW; D26CD1E63AAE0EB9 CRC64;
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             Miyazaki T., Miyazaki J., Nishiyama M., Yamane H.;
"Characterization of a LysN, the 4th enzyme in lysine biosynthesis, in an extremely thermophilic bacterium, Thermus thermophilus HB27.";
Submitted (NOV-2002) to the EMBL/GenBank/DDBJ databases.
EMBL, AB097117, BA76940.1;
"Myochetical protein.
                                                                                                                                                                                                                                                                                                                                                                           Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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Bukaryota, Viridiplantea, Streptophyta, Embryophyta, Tracheophyta;
Spermatophyta, Magnoliophyta, eudicocyledons; core eudicots; rosids;
eurosids 1; Fabales, Fabaceae, Papilionoldeae, Phaseoleae;
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Hystricognathi; Caviidae.
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Pred. No. 5.6e+03;
2; Mismatches 0; Indels
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01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-UJN-2002 (TrEMBLrel. 21, Last annotation update)
GP-C83=CORTICOSTATIC peptide (Fragment).
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Biochem. Biophys. Res. Commun. 180:558-565(1991)
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Phytochemistry 31:731-735(1992).
PIR; C61497; C61497.
                                                                                                                                                                                                                                                                                                             25.6%;
60.0%;
                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 60.v-
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SEQUENCE FROM N.A.
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NCBI_TaxID=3891;
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MEDILINE=9635021; PubMed=8757037;
Mayer S.A., Rubin B.S., Starman B.J., Byers P.H.;
"Spontaneous multivessel cervical artery dissection in a patient with a substitution of alanine for glycine (G13A) in the alpha 1 (I) chain
                                                                                                                                                                                                                                                                                                                                                        MEDLINE=94059921; PubMed=8241100;
Novotny W.F., Maffi T., Mehta R.L., Milner P.G.;
Nototny W.F., Maffi T., Mehta R.L., Milner P.G.;
"Identification of novel heparin-releasable proteins, as well as the cytokines midkine and pleiotrophin, in human postheparin plasma.";
Arterioscler. Thromb. 13:1798-1805(1993).
SEQUENCE 15 AA; 1454 MW; 0585FEF4FE8F6265 CRC64;
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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15 15
15 AA; 1413 MW; 4CABD720AAF780FC CRC64;
OUCC2 PKELLEL 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-MAY-2000 (TrEMBLrel. 14, Last annotation update)
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01-JUL-1997 (TrEMBLrel. 04, Last sequence update)
01-JUL-2003 (TrEMBLrel. 24, Last annotation update)
Type I collagen alpha 1(I) chain (Fragment).
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Neurology 47:552-556(1996).
EMBL; S83315; AAB50780.2; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3; Conservative
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Best Local Similarity
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5 PVPIPL 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 9 PRPLP 13
                                                                                                                                                                                                                                                                          NCBI_TaxID=9606;
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Fri Apr
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|| || 5 PRGLP 9

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MEDLINE=94320083; PubMed=8044796; MSDLINE=94320083; PubMed=8044796; Magalla S.R., Spindel E.R.; Wegalla S.R., Spindel E.R.; Functional analysis of the 5'-flanking region of the human gastrin-releasing peptide gene in small cell lung carcinoma cell lines."; Cancer Res. 54:4461-4467(1994).

EMBL; $73265; Amb14116.1; -. GO; GO:0005634; C:nucleus; NAS.
GO; GO:0006355; P:regulation of transcription, DNA-dependent; NAS. SRQUENCE 9 AA; 1070 MW; 77FE37672B040864 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          noss.
Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                            Q16220;
01-NOV-1996 (TrEMBLrel. 01, Created)
01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
HGRP protein (Fragment).
                                                                                                                     9 AA.
                                                                                                                     PRT;
                                                                                                               PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NCBI TaxID=9606;
                                                                                                          Q16220
RESULT 15
016220
D1 016220
DT 01-NOV
```

1; Indels Query Match

24.4%; Score 20; DB 4; Length 9;
Best Local Similarity 66.7%; Pred. No. 1e+06;
Matches 4; Conservative 1; Mismatches 1; Indels 10 RPLPVV 15 4 RELPLV 9 qq

Search completed: April 2, 2004, 10:11:07 Job time : 32.3333 secs

0

Gaps

0;

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GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
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OM protein - protein search, using sw model

April 2, 2004, 10:06:43 ; Search time 14 Seconds (without alignments) 55.314 Million cell updates/sec Run on:

US-10-009-049-3 82 1 CRAHSEVSPRPLPVV 15 Perfect score:

BLOSUM62 Scoring table: Sequence:

389414 seqs, 51625971 residues Searched:

Gapop 10.0 , Gapext 0.5

Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 15

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Issued_Patents_AA:*
1: /cgn2_6/ptodata/2/iaa/5A_CCMB.pep:*
2: /cgn2_6/ptodata/2/iaa/5A_CCMB.pep:*
3: /cgn2_6/ptodata/2/iaa/6A_CCMB.pep:*
4: /cgn2_6/ptodata/2/iaa/6B_CCMB.pep:*
5: /cgn2_6/ptodata/2/iaa/PcTUS_COMB.pep:*
6: /cgn2_6/ptodata/2/iaa/PcTUS_COMB.pep:* Database :

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

		Appl	Appl	App1	Appl	Appl	ppli	Appl ,	Appl	Appl	App1	App	App	App		App					App	Appl	Appl	Appl	Appl	Appl	Appl	Appl
	렱	30,			92,		3, 8			29,		167,	170,	167,	167,	219,	255,	255,	783,	410,	410,	86,	86,	86,				67,
	Description	Seguence	Seguence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence								Sequence	Sequence	Sequence	Seguence	Sequence	Sequence	Sequence
SOUTHERNESS	ID	US-08-230-047-30	US-08-230-047-39	US-08-602-999A-92	- 1	-124-9	-60-	-08-336	US-08-652-877-29	US-08-476-515A-29	-09-845-	US-08-190-788A-167	US-08-383-474B-170	US-08-465-391A-167	-08-46	-463-	US-08-602-999A-255	US-09-500-124-255	US-09-428-082B-783	-08-602-9	-09-500-124-41		US-08-278-865-86	-09-500-12	-08-60	US-08-278-865-62	-005-60-	US-08-602-999A-67
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	Length	15	13	13	13	13	6	11	11	11	11	12	12	12	12	12	12	12	12	15	15	13	13	13	14	14	14	14
₩	Query Match	47.6	42.7	42.7	42.7	42.7	41.5	41.5	41.5	41.5	41.5	41.5	41.5	41.5	41.5			41.5						ö	0	•	40.2	37.8
	Score	39	35	35	35	35	34	34	34	34	34	34	34	34	34	34	34	34	34	34	34	33	33	33	33	33	33	31
	Result No.	н	7	м	4	Ŋ	w	7	80	6	10	11	12	13	14	15	16	17	18	19	50	21	22	23	24	25	56	27

Sequence 67, Appl		Sequence 29, Appl	96	355,		Sequence 152, App	355,		24, A	34,		34,		Sequence 218, App	221, A	Sequence 218, App	Sequence 218, App
US-08-278-865-67	US-09-500-124-67	US-08-230-047-29	US-08-630-916A-96	US-08-602-999A-355	US-08-602-999A-396	US-08-630-915A-152	US-09-500-124-355	US-09-500-124-396	US-08-336-343A-24	US-08-652-877-34	US-08-652-877-36	US-08-476-515A-34	US-08-476-515A-36	US-08-190-788A-218	US-08-383-474B-221	US-08-465-391A-218	US-08-464-538B-218
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14	14	15	15	15	15	15	15	15	11	11	11	11	11	12	12	12	12
37.8	37.8	37.8	37.8	37.8	37.8	37.8	37.8	37.8	36.6	36.6	36.6	36.6	36.6	36.6	36.6	36.6	36.6
31	31	31	31	31	31	31	31	31	30	30	30	30	30	30	30	30	30
28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45

ALIGNMENTS

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Sequence 30.
Seque
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MEDIUM TYPE: Floppy disk
COMPUTER: Macintosh
OPERATING SYSTEM: System 7.1
SOFTWARE: Word 5.0 (Patentin)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/230,047
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION: 435
ATTORNEY, AGENT INFORMATION:
NAME: GOOGMAAN, ROSANNE
REGISTRATION NUMBER: 32,534
REFERENCE/DOCKET NUMBER: A1465-US
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: (610) 454-3808
INFORMATION FOR SEQ ID NO: 30:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (610) 454-3817
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        19-APR-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19426
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEPHONE:
US-08-230-047-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-08-230-047-30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGIH:
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0 0; Gaps 47.6%; Score 39; DB 1; Length 15; 70.0%; Pred. No. 3.3; tive 0; Mismatches 3; Indels Query Match
Best Local Similarity 70.0°
Matches 7; Conservative 4 HSFVSPRPLP 13 ઇ

1 HSHFHPRPLP 10

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TELECOMMUNICATION INFORMATION:
TELEPHONE: (703) 413-3000
TELEFAX: (703) 413-220
TELEFAX: 248855 OPAT UR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE:
CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                                                                                                                               unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      8 SPRPLPW 15
                                                                                                                                                                                                                                                                                                                                                                            amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Virginia
: U.S.A.
                                                                                                                FILING DATE: 16 CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE:
ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                   US-08-602-999A-92
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 4
US-08-278-865-92
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                                                   GENERAL INCORNATION:
APPLICANT: Searfoss III, George H.
APPLICANT: Ivashchenko, Yuri D.
APPLICANT: Jaye, Michael C.
TITLE OF INVENTION: BENESSION CLONING OF C-SRC SH3 BINDING TITLE OF INVENTION: PROTEIN
NUMBER OF SEQUENCES: 40
CORRESPONDENCE ADDRESS:
ADDRESSES: Rhone-Poulenc Rorer Inc.
STREET: 500 Arcola Road, 33C43
CITY: Collegeville
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 35; DB 1; Length 13; Pred. No. 13; 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: RIDER, James E.
TITLE OF INVENTION: SH3 BINDING PEPTIDES AND METHODS OF
TITLE OF INVENTION: ISOLATING AND USING SAME
NUMBER OF SEQUENCES: 467
CORRESPONDENCE ADDRESS:
ADDRESSEE: Pennie & Edmonds
STREET: 1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: Macintosh OPERATING SYSTEM 7.1 SOFTWARE: GOTTWARE: WORD 5.0 (Patentin) CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/230,047 FILING DATE: 19-APR-1994 CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
APPLICANT: SPARKS, Andrew B.
APPLICANT: RAY, Brian K.
APPLICANT: THORN, Judith M.
APPLICANT: OUILLIAM, Lawrence A.
APPLICANT: PDER, Channing J.
APPLICANT: RIDER, James B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 92, Application US/08602999A
Patent No. 6184205
                    Sequence 39, Application US/08230047
Patent No. 5541109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
NAME: Goodman, Rosanne
REGIESTRATION NUMBER: 32,534
REFERENCE/DOCKET NUMBER: A146;
TELECOMMUNICATION INFORMATION:
TELEPHONE: (610) 454-3817
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: (610) 454-3808
INFORMATION FOR SEQ 1D NO: 39:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            42.7%;
                                                                                                                                                                                                                                                                                                                                                      E: Floppy disk
Macintosh
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             13 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 70.0.
7, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY: linear
MOLECULE TYPE: peptide
US-08-230-047-39
                                                                                                                                                                                                                                                                                                             ZIP: 19426
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          4 HSFVSPRPLP 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               3 HSAVPPRPGP 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STATE: New York COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET: 1155 Av CITY: New York
                                                                                                                                                                                                                                                                                              USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 3
US-08-602-999A-92
US-08-230-047-39
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                                                                                                                                                                                                                                                                          STATE:
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WESTITY PREASONED FROM THE COMPUTER IN TYPE FLORY disk COMPUTER. HEADER PREASONED FROM MILE COMPUTER. HEADER COMPUTER. HEADER
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TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 869-9741/8864
TELEX: (6141 PENNIE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; LOCATION: (1)...(9)
; OTHER INFORMATION: Synthetic Peptide
US-09-086-168B-3
                                                                                                                                                                                                                                                                                                                                                                                                                                FEATURE:
OTHER INFORMATION: SH3 test peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 29, Application US/08336343A Patent No. 5677144 GENERAL INFORMATION:
                                                                                               Sequence 3, Application US/09086168B; Patent No. 6638941
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            E: Floppy disk
IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Coruzzi, Laura A. REGISTRATION NUMBER: 30,742
                                                                                                                                                                                                                                                                                                                                                                                          TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: New York
STATE: New York
COUNTRY: U.S.A.
ZIP: 10036-2711
COMPUTER READABLE FORM:
MEDIUM TYPE: Ploppy di
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME/KEY: MISC_FEATURE
      2 TPRPLPLI 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             9 PRPLPV 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 PRPLPV 6
                                                                                                                                               GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-08-336-343A-29
                                                                                  US-09-086-168B-3
                                                                                                                                                                                                                                                                                                                                                  SEQ ID NO 3
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42.7%; Score 35; DB 4; Length 13;
Best Local Similarity 62.5%; Pred. No. 13;
Matches 5; Conservative 3; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: KAY, Brian K.
APPLICANT: THORN, Judith M.
APPLICANT: OUILLIAM, Lawrence A.
APPLICANT: DER, Channing J.
APPLICANT: ROWIKES, Dana M.
APPLICANT: RIDER, James E.
TITLE OF INVENTION: SH3 BINDING PEPTIDES AND METHODS OF
                                                                                                                                                            42.7%; Score 35; DB 4; Length 13; 62.5%; Pred. No. 13; tive 3; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/500,124
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
APPLICATION NUMBER:
APPLICATION TO THE TENDER OF THING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Misrock, S. Leslie
REGISTRATION NUMBER: 18,872
REFERENCE/DOCKET NUMBER: 1101-202
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES: 467
CORRESPONDENCE ADDRESSS:
ADDRESSE: Pennie & Edmonds
STREET: 1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY:
ZIF: 10036-2711
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IRM PC compatible
COMPUTER: READABLE
COMPUTER: ISM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                      Sequence 92, Application US/09500124
Patent No. 6432920
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 790-9090
TELEFAX: (212) 869-9741/8864
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 92:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                 SPARKS, Andrew B.
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 13 amino acids TYPE: amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 13 amino acids
                                                                                                                                                                                 Best Local Similarity 62.5
Matches 5, Conservative
                                                                       ; TOPOLOGY: unknown; MOLECULE TYPE: peptide US-08-278-865-92
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    unknown
                                                                                                                                                                                                                                            8 SPRPLPVV 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         8 SPRPLPWV 15
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2 TPRPLPLI 9
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APPLICANT: SPARKS
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                                                                                                                                                                                                                                                                                                                                             RESULT 5
US-09-500-124-92
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                                                                                                                                                              Query Match
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APPLICANT: Castelhano, Arlindo
APPLICANT: Mitter, David
TITLE OF INVENTION: CONFORMATIONALLY CONSTRAINED PEPTIDOMIMETICS AS BETA-TURN
TITLE OF INVENTION: TEMPLATES AND MODULATORS OF SH3 DOMAINS
FILE REFERENCE: 60384-B
CURRENT APPLICATION NUMBER: US/09/086,168B
CURRENT FILING DATE: 1998-05-28
NUMBER OF SEQ ID NOS: 5
SOFTWARE: PATENTIN version 3.1
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APPLICANT: Alves, Frauke
APPLICANT: Alves, Frauke
TITLE OF INVENTION:
CCK-2, A No. 5677144el Receptor Tyrosine Kinase
NUMBER OF SEQUENCES: 43
CORRESPONDENCE ADDRESS:
ADDRESSEE: Pennie & Edmonds
STREET: 1155 Avenue of the Americas
CITY: New York
CITY: New York
COUNTRY: U.S.A.
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ATTORNEY/ALPANO... 435
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INFORMATION FOR SEQ ID NO: 29:
SEQUENCE CHARACTERISTICS:
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US-08-476-515A-29
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Patent No. 6187548

GENERAL INFORMATION:
APPLICANT: Akerstrom, Goran
APPLICANT: Rask, Lars
APPLICANT: Crumley, Gregg R.
APPLICANT: Murray, Gregg R.
APPLICANT: Mirray, Gregg R.
APPLICANT: Hjalm, Goran
TILLE OF INVENTION: Human Calcium Sensor Protein, Fragments
TILLE OF INVENTION: Thereof and DNA Encoding Same
NUMBER OF SEQUENCES: 106
CORRESPONDENCE ADDRESS:
                                                                                                                 Query Match
41.5%; Score 34; DB 1; Length 11;
Best Local Similarity 100.0%; Pred. No. 16;
Matches 6; Conservative 0; Mismatches 0; Indels
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REGISTRATION NUMBER: 29,699
REPRENCE/DOCKET NUMBER: 41355E-US
TELECOMMUNICATION INFORMATION:
TELEPHONE: 610-454-3808
INFORMATION FOR SEQ ID NO: 29:
SEQUENCE CHARACTERISTICS:
LENGTH: 11 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: DCT/US95/15203
FILING DATE: 22-NOV-1995
PRIOR APPLICATION NUMBER: US 08/344,836
FILING DATE: 23-NOV-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/344,836
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/487,314
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEE: Rhone-Poulenc Rorer Inc.: 500 Arcola Rd., 3C43
Collegeville
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OPERATING SYSTEM: System 7.5.1
SOFTWARE: Word 6.0 (Patentin)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/652,877
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE: 07-JUNE-1995
ATTORNEY/AGENT INFORMATION:
NAME: Savitzky, Martin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY: USA
ZIP: 19426-0107
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Macintosh
LENGTH: 11 amino acids
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            ; TYPE: amino acid
; TOPOLOGY: unknown
; MOLECULE TYPE: peptide
US-08-336-343A-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE: peptide
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                                                                                                                                                                                                                                                                                                APPLICANT: Akerstrom, Goran
APPLICANT: Juhlin, Claes
APPLICANT: ALL Claes
APPLICANT: Rare, Carence C.
APPLICANT: Morse, Clarence C.
APPLICANT: Milm, Goran
APPLICANT: Hialm, Goran
TITLE OF INVENTION: Thereof and DNA Encoding Same
NUMBER OF EQUENCES: 84
CORRESPONDENCE ADDRESS:
ADDRESSEE: Martin Savitzky
STREET: Rhone-Poulenc Rorer Inc., 500 Arcola Rd.;
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Best Local Similarity 100.0%; Pred. No. 16;
Matches 6; Conservative 0; Mismatches 0; Indels
Query Match
Best Local Similarity 100.0%; Pred. No. 16;
Matches 6; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: Compad PC
OPERATING SYSTEM: Windows 95
SOFTWARE: Word 7.0 (Patentin)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/476,515A
FILING DATE: 07-JUN-1995
CLASSIFICATION DATA:
APPLICATION NUMBER: US 08/344,836
FILING DATE: 23-NOV-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/344,836
FILING DATE: 24-MAY-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: WO PCT/SE94/00483
FILING DATE: 24-MAY-1993
ATTOMNEY/ABSHI INFORMATION:
                                                                                                                                                                                                                                       Sequence 29, Application US/08476515A Patent No. 6239270
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Savitzky, Martin
REGISTRATION UNDRER: 29,699
REFERENCE/DOCKET NUMBER: A135:
TELECOMMUNICATION INFORMATION:
TELEPANE: 610-454-3816
TELEPAX: 610-454-3808
INFORMATION FOR SEQ ID NO: 29:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ZIP: 19426-0107
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 11 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               HYPOTHETICAL: NO FRAGMENT TYPE: internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MOLECULE TYPE: peptide HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STREET: 3C43,
CITY: Collegeville
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: amino acid STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             linear
                                                                                         9 PRPLPV 14
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INFORMATION FOR SEQ ID NO: 167:
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US-08-383-474B-170
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                                                                                                                               Sequence 17, Application US/09845917A

Patent No. 6653529

GENERAL INPORMATION

GENERAL INPORMATION

APPLICANT: Bogaert, Thierry

APPLICANT: Vandekerckhove, Joel

TITLE OF INVENTION: PROCESES FOR THE IDENTIFICATION OF COMPOUNDS WHICH

TITLE OF INVENTION: OWNTROL CELL BEHAVIOUR, THE COMPOUNDS IDENTIFIED AND

TITLE OF INVENTION: USE IN THE CONTROL OF CELL BEHAVIOUR

TITLE OF INVENTION: USE IN THE CONTROL OF CELL BEHAVIOUR

TITLE OF INVENTION: USE IN THE CONTROL OF CELL BEHAVIOUR

CURRENT APPLICATION NUMBER: US/09/845,917A

CURRENT FILING DATE: 2011-04-30

NUMBER OF SEQ ID NOS: 58
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| Patent No. 5608035
| CENERAL INFORMATION: Sequence 167, Application US/08190788A
| CENERAL INFORMATION: APPLICANT: Vanofsky, Stephen D. APPLICANT: Bardwin, David W. APPLICANT: Bardwin, David W. APPLICANT: Jacobs, Jeff W. TITLE OF INVENTION: Peptides and Compounds That Bind to the TITLE OF INVENTION: IL-1 Receptor NUMBER OF SEQUENCES: 312
| CORRESPONDENCE ADDRESS: ADDRESSEE: Affymax Technologies N.V.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 41.5%; Score 34; DB 4; Length 11; 100.0%; Pred. No. 16; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDIUM TYPE: Floppy disk COMPUTER: IBM PC COMPATIBLE COMPUTER: IBM PC COMPATIBLE COMPATING SYSTEM: PC-DOS/MS-DOS SOFWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/190,788A
FILING DATE: 02-FEB-1994
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/847,567
FILING DATE: 05-MAR-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: PRT
ORGANISM: Caenorhabditis elegans
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STREET: 4001 Miranda Avenue
CITY: Palo Alto
STATE: California
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Stevens, Lauren L.
REGISTRATION NUMBER: 36,691
REPERENCE/DOCKET NUMBER: 10
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-496-2300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity 100.0
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PRPLPV 7
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Fatent No. 5767234
GENERAL INFORMATION:
APPLICANT: Manofsky, Stephen D.
APPLICANT: Barrett, Ronald W.
APPLICANT: Baldwin, David N.
APPLICANT: Baldwin, David N.
APPLICANT: Baldwin, David N.
APPLICANT: Beldwin, David N.
APPLICANT: Horby 100: The IL-1 Receptor
NUMBER OF SEQUENCES: 146
CORRESPONDENCE ADDRESS:
                                                                                                                                                                        41.5%; Score 34; DB 1; Length 12; 54.5%; Pred. No. 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  41.5%; Score 34; DB 1; Length 12; 54.5%; Pred. No. 17; 17; ive 1; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATIOS SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/383,474B
FILING DATE: 01-FEB-1995
CLASSIFICATION: 530
PRIOR APPLICATION: 530
PRIOR APPLICATION NUMBER: US 08/190,788
FILING DATE: 02-FEB-1994
CLASSIFICATION: 530
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE: Townsend & Townsend & Crew LLP STREET: Two Embarcadero Center, 8th Floor CITY: San Francisco STATE: Callfornia
                                                                                                                                                                                                                  1; Mismatches
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NAME: Stevens, Lauren L.
REGISTRATION NUMBER: 36,691
REFERENCE/DOCKET NUMBER: 101
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: 415-496-2300
TELEFAX: 415-424-0832
SEQUENCE CHARACTERISTICS:
LENGTH: 12 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELBFAX: 415-424-0832
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  12 amino acids
                                                                                                                                                                                                                  6; Conservative
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                                                                  single
                                                                                                     MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                            4 HSFVSPRPLPV 14
                                                                                                                                                                                                                                                                                                    2 HQFVQPYALPL 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4 HSFVSPRPLPV 14
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                                          TYPE: amino acid
STRANDEDNESS: Bir
TOPOLOGY: linear
                                                                                                                                                                          Query Match
Best Local Similarity
Matches 6; Conserv
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Best Local Similarity
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                                                                                        APPLICANT: Barrett, Ronald W.
APPLICANT: Yanofsky, Stephen D.
APPLICANT: Yanofsky, Stephen D.
APPLICANT: Baldwin, David
APPLICANT: Jacobs, Jeff W.
APPLICANT: Leahy, Phillipe R.
APPLICANT: Pottorf, Richard S.
TITLE OF INVENTION: Peptides and Compounds That Bind to the TITLE OF INVENTION: IL-1 Receptor
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
41.5%; Score 34; DB 1; Length 12;
Best Local Similarity 54.5%; Pred. No. 17;
Matches 6; Conservative 1; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                               One Market Plaza, Steuart Tower, Suite 2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION DATA:

SOFTWARE: Patentin Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/465,391A

FILING DATE: 05-JUN-1995

CLASSIFICATION: 514

PRIOR APPLICATION UNBER: US 08/373,474

FILING DATE: 01-FEB.1995

CLASSIFICATION NUMBER: US 08/130,788

FILING DATE: 02-FEB-1994

CLASSIFICATION: 514

ATTORNEY/AGENT INFORMATION:

NAME: NO. 5786331viel, Vern

NAME: NO. 5786331viel, Vern

NAME: NO. 5786331viel, Vern

NAME: NO. 5786331viel, Vern

NAME: NAME: NUMBER: 32,483

REGESTRATION NUMBER: 32,483

REGESTRATION NUMBER: 32,483
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                                                                                                                                                                                                                                                                                                                                                                   Townsend and Townsend and Crew
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Patent No. 5861476
GENERAL INFORMATION:
APPLICANT: Barrett, Ronald W.
APPLICANT: Yanofsky, Stephen D.
US-08-465-391A-167
; Sequence 167, Application US/08465391A
; Patent No. 5786331
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER: 16
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-326-2400
TELEFAX: 415-326-2422
INFORMATION FOR SEQ ID NO: 167:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         12 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            4 HSFVSPRPLPV 14
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                                                                                                                                                                                                                                                                                                                                                                                                                   CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 linear
                                                                              GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH
                                                                                                                                                                                                                                                                                                                                                                                            STREET:
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Gaps
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| Patent No. 5880096
| GENERAL INFORMATION:
| APPLICANT: Barrett, Ronald W. APPLICANT: Barrett, Ronald W. APPLICANT: Barrett, Ronald W. TITLE OF INVENTION: IL-1 Receptor NUMBER OF SEQUENCES: 392
| CORRESPONDENCE ADDRESS: ADDRESSE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco CITY: Eighth Floor CORMENTED COMMUNES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ;
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APPLICANT: Baldwin, David
APPLICANT: Jacobs, Jeff W.
APPLICANT: Bovy, Phillipe R.
APPLICANT: Leahy, Ellen M.
APPLICANT: Pottorf, Richard S.
TITLE OF INVENTION: Peptides and Compounds That Bind to the NINUMBER OF SEQUENCES: 402
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 34; DB 2; Length 12;
Pred. No. 17;
1; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION DATA:

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/464,538B
FILING DATE: 05-UNW-1995
CLASSIFICATION ATA:

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US/08/464,538B
FILING DATE: 05-UNW-1995
CLASSIFICATION DATA:

APPLICATION NUMBER: US 08/373,474
FILING DATE: 01-FEB-1995
CLASSIFICATION SATA:

APPLICATION NUMBER: US 08/190,788
FILING DATE: 02-FEB-1994
CLASSIFICATION SATA:

ATTORNEY/AGENT INFORMATION:

NAME: Smith, William M.

REGISTRATION NUMBER: 16528A-001810
TELECOMMUNICATION NUMBER: 16528A-001810
TELEPHONE: 415-326-2442
INFORMATION FOR SEC 1D NO: 167:
                                                                                                                                                                                                  Townsend and Townsend and Crew LLP
                                                                                                                                                                                                                         Two Embarcadero Center, 8th Floor
                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                41.5%;
54.5%;
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Best Local Similarity 54.0.,
Best Local 6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              12 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; TOPOLOGY: linear; MOLECULE TYPE: peptide US-08-464-538B-167
                                                                                                                                                                             CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4 HSFVSPRPLPV 14
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                                                                                                                                                                                                                         STREET: Two Embarcad
CITY: San Francisco
STATE: California
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                                                                                                                                                                                                ADDRESSEE:
                                                                                                                                                                                                                                                                                    COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 15
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Query Match
Best Local Similarity 54.5%; Pred. No. 17;
Matches 6; Conservative 1; Mismatches 4; Indels
COMPUTER READBLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IEMP FC compatible
COMPUTER: IEMP FC compatible
COMPUTER: IEMP FC compatible
COMPUTER: IEMP FC compatible
COMPUTER: Datentin Release #1.0, Version #1.30
CURRENTE APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US/08/463,076E
FILING DATE: 05-UN-1995
CLASSIFICATION: 104
ATTORNEY/AGENT INFORMATION:
REGISTRATION NUMBER: 39,381
REFERENCE/DOCKET NUMBER: 16528A-001850US
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECHONE: (415) 576-0200
TELECHONE: (415) 576-0200
TELECHONE: (415) 576-0300
INFORMATION FOR SEQ ID NO: 219:
SEQUENCE CHARACTERISTICS:
LENGTH: 12 amino acids
TYPE: amino acid
STYPE: Innear
MOLECULE TYPE: peptide
US-08-463-076E-219
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0; Gaps

Search completed: April 2, 2004, 10:12:45 Job time : 14 secs

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April 2, 2004, 10:11:14 ; Search time 32.3333 Seconds (without alignments) 121.720 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                               GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
                                                                                                                                                                                                                                    Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                               1070241 seqs, 262374223 residues
                                                                                                                                                                                                                                                                                                  Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                  OM protein - protein search, using sw model
                                                                                                                                                                          BLOSUM62
Gapop 10.0 , Gapext 0.5
                                                                                                                                                 1 CRAHSFVSPRPLPVV 15
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                                                                                                                                                                                                                                                               Minimum DB seq length: 0
Maximum DB seq length: 15
                                                                                                                        Title:
Perfect score:
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                                                                                                                                                  Sequence:
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

	ריידים פרן אניידים פרן	Desci Trans	Sequence 92, Appl	Sequence 92, Appl	Sequence 17, Appl	Sequence 17, Appl	Sequence 783, App	Sequence 783, App	Sequence 783, App	Sequence 255, App		Sequence 22, Appl	Sequence 86, Appl	Sequence 86, Appl	Sequence 62, Appl	Sequence 62, Appl	Sequence 67, Appl
	£		US-09-938-315-92	US-10-161-791-92	US-09-945-917-17	US-09-845-917A-17	US-10-609-217-783	US-10-632-388-783	US-10-651-723-783	US-10-161-791-255	US-10-161-791-410	US-09-999-686-22	US-09-938-315-86	US-10-161-791-86	US-09-938-315-62	US-10-161-791-62	US-09-938-315-67
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Sequence 67, Appl Sequence 152, Appl Sequence 96, Appl Sequence 355, App	4 4 8 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6	Sequence 835, App Sequence 835, App Sequence 30, Appl Sequence 281, Appl Sequence 85, Appl	8 8 8 1 8 1 9 1 9 1 9 1 9 1 9 1 9 1 9 1	Sequence 14, Appl Sequence 14, Appl Sequence 14, Appl Sequence 960, App Sequence 960, App	96 68 30 114 26 27 28
14 US-10-161-791-67 9 US-09-879-957-152 14 US-10-185-050-96 14 US-10-161-791-355	US-10-161-71-3 US-09-572-404B- US-10-149-13B-4 US-10-609-217-8	US-10 US-10 US-10 US-10	4 US-10-161 4 US-10-161 2 US-10-398 4 US-10-158	14 US-10.157-7789-14 14 US-10-254-446A-14 14 US-10-155-883B-14 9 US-09-895-814-960 13 US-10-012-896-960	14 US-10-294-025-960 14 US-10-232-544-68 14 US-10-161-791-306 9 US-09-815-837-114 14 US-10-008-524A-26 14 US-10-008-524A-27 14 US-10-008-524A-27
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ALIGNMENTS

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US-09-388-115-92
Sequence 92, Application US/0938315
Patent No. US20020010085A1
Sequence 92, Application US/0938315
Patent No. US20020010085A1
SPARKS, ANDREW B.
THORNATION:
SPARKS, ANDREW B.
THORNATION:
CULLIAM, LAWRENCE A.
DER, CHANNING W.
SPINCE SH3 BINDING PEPTIDES AND METHODS OF THORNATION: SCC SH3 BINDING PEPTIDES AND METHODS OF CORRESPONDENCE: OBLON, SPIVAK, MCCLELLAND, MAIER & NEUSTADT, STREET: 175 S.
STREET: 175 S. Jefferson Davis Highway, Suite 400 CITY: Arlington
STRATE: Virginia
COUNTRY: U.S.A.
ZITY: Arlington
STATE: Propry disk
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM C compatible
OPERATING SYSTEM: PC-DS5/MS-DS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT SASTEM: PC-DS6/MS-DS
SOFTWARE: PatentIn Release #1.0, Version #1.25
FILING DATE: 23-Aug-2001
CLASSIFICATION NUMBER: 19,398
TELESCHANNICATION INFORMATION:
REFERENCE/DOCKET NUMBER: 49,908-007-0
TELEPRAN: (703) 413-2220
TELESCHAUNICATION INFORMATION:
TELEPRAN: (703) 413-2220
TELEEX: 248855 OPAT UR
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                                                                                                                                                                                    42.7%; Score 35; DB 9; Length 13; 62.5%; Pred. No. 1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: QUILLIAM, LAWRENCE A.
APPLICANT: DER, Channing J.
APPLICANT: FOWLKES, Dana M.
APPLICANT: FOWLKES, Dana M.
APPLICANT: RIDER, James E.
TITLE OF INVENTION: SH3 BINDING PEPTIDES AND METHODS OF
NUMBER OF SEQUENCES: 467
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                               0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/NS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/161,791
                                                                                                                                                                                                                               3; Mismatches
                                                                                  TOPOLOGY: unknown
MOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: SEQ ID NO: 92:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE: Pennie & Edmonds
STREET: 1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/602,999
FILING DATE: 16-FEB-1996
ATTORNEY/AGENT INFORMATION:
NAME: Misrock, S. Leslie
REGISTRATION NUMBER: 18,872
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1101-202
                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 92, Application US/10161791
Publication No. US20030186863A1
GENERAL INFORMATION:
INFORMATION FOR SEQ ID NO: 92:
SEQUENCE CHARACTERISTICS:
LENGTH: 13 amino acidd
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REGISTRATION NUMBER: 18,872
REFERENCE/DOCKET NUMBER: 1101
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 790-9090
TELEFAX: (212) 869-9741/8864
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: SPARKS, Andrew B. APPLICANT: KAY, Brian K. APPLICANT: THORN, Judith M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 92:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: amino acide TOPOLOGY: unto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity 62.5.
                                                                                                                                                                                                                             5; Conservative
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LENGTH: 13 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE: peptide US-10-161-791-92
                                                                                                                                                                                                                                                                     8 SPRPLPVV 15
                                                                                                                                                                                                                                                                                                           2 TPRPLPLI 9
                                                                                                                                                                                                     Best Local Similarity
Matches 5; Conserv
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: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     10036-2711
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                          US-10-161-791-92
                                                                                                                                            US-09-938-315-92
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY:
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8 SPRPLPVV 15

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                                                                                                        US-09-945-917-17

US-09-945-917-17

Sequence 17, Application US/09945917

Sequence 17, Application US/09945917

Publication No. US20030042381A1

SEQUENCE 17: APPLICAMON.

APPLICAMON.

APPLICAMON.

TITLE OF INVENTION: CONTROL CELL BEHAVIOUR, THE COMPOUNDS WHICH

TITLE OF INVENTION: PHARMACEUTICAL COMPOSITIONS CONTIANING THEM AND THEIR

FILE OF INVENTION: UNBER: US-09-945, 917

CURRENT APPLICATION NUMBER: US/09/945, 917

CURRENT FILING DATE: 1998-09-21

NUMBER OF SEQ ID NOS: 58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Bogaert, Thierry
APPLICANT: Vandekerckhove, Joel
TITLE OF INVENTION: PROCESSES FOR THE IDENTIFICATION OF COMPOUNDS WHICH
TITLE OF INVENTION: CONTROL CELL BEHAVIOUR, THE COMPOUNDS IDENTIFIED AND
TITLE OF INVENTION: PHARMACEUTICAL COMPOSITIONS CONTIANING THEM AND THEIR
FILLE REPERENCE: P/14-1
CURRENT APPLICATION NUMBER: US/09/845,917A
CURRENT FILING DATE: 2001-04-30
NUMBER OF SEQ ID NOS: 58
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41.5%; Score 34; DB 10; L.
Best Local Similarity 100.0%; Pred. No. 1.2e+02;
Matches 6; Conservative 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Caenorhabditis elegans
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT
ORGANISM: Caenorhabditis elegans
                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 17
LENGTH: 11
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Best Local Similarity 100.0
Matches 6; Conservative
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2 TPRPLPLI 9
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
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REGISTRATION NUMBER: 18,872
REFERENCE/DOCKET NUMBER: 110
TELECOMMUNICATION:
TELEPHONE: (212) 790-9090
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 255:
                                                                                                                                                                                                                                                                                   TYPE: PRT ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 12 amino acids
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CITY: New York
STATE: New York
COUNTRY: U.S.A.
ZIP: 10036-2711
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM PC
OPERATING SYSTEM:
SOFTWARE: PAtentI
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CLASSIFICATION:
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Publication No. US20040053845A1
GENERAL INFORMATION:
APPLICANT: ETGE, ULRICH
APPLICANT: LIU CHUAN-FA
APPLICANT: LIU CHUAN-FA
APPLICANT: GENETHAM, JANET C.
TILLE OF INVENTION: MODIFIED PEPTIDES AS THERAPEUTIC AGENTS
TILLE OF INVENTION: MODIFIED PEPTIDES AS THERAPEUTIC AGENTS
CURRENT FILING DATE: 2003-07-31
PRIOR APPLICATION NUMBER: US/10/632,388
CURRENT FILING DATE: 1999-10-22
PRIOR FILING DATE: 1999-10-23
PRIOR FILING DATE: 1998-10-23
PRIOR FILING DATE: 1998-10-23
NUMBER OF SEQ ID NOS: 1133
SOFTWARE PATENTIN VERSION 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                              41.5%; Score 34; DB 12; Length 12; 54.5%; Pred. No. 1.4e+02; tive 1; Mismatches 4; Indels
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                 APPLICANT: BOONE, THOMAS CHARLES
TITLE OF INVENTION: MODIFIED PEPTIDES AS THERAPEUTIC AGENTS
                                                        FILE REFERENCE: A-527
CURRENT APPLICATION NUMBER: US/10/609,217
FRICH APPLICATION NUMBER: US/09/428,0828
FRICH APPLICATION NUMBER: US/09/428,0828
FRICH FILING DATE: 1999-10-22
FRICH FILING DATE: 1998-10-23
NUMBER OF SEQ ID NOS: 1133
SOFTWARE: Patentin Version 3.1
LENGTH: 12
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) OTHER INFORMATION: IL-1 ANTAGONIST PEPTIDE US-10-609-217-783
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Publication No. US2004057953A1
GENERAL INFORMATION:
APPLICANT: FEIGE, ULRICH
APPLICANT: LIU, CHUAN-FA
APPLICANT: CHEETHAM, JANET C.
                                                                                                                                                                                                                                                                                                 TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: PRT
ORGANISM: Artificial Sequence
CHEETHAM, JANET C.
                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 54.5'
Matches 6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity 54.5
Matches 6, Conservative
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APPLICANT:
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Gaps
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APPLICANT: KAY, Brian K.
APPLICANT: THORN, Judith M.
APPLICANT: THORN, Judith M.
APPLICANT: OUTLIAM, Lawrence A.
APPLICANT: DER, Channing J.
APPLICANT: POMLKES, Dana M.
APPLICANT: RIDER, James E.
TITLE OF INVENTION: SRJ BINDING PEPTIDES AND METHODS OF TITLE OF INVENTION: ISOLATING AND USING SAME
NUMBER OF SEQUENCES: 467
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 12;
APPLICANT: BOONE, THOMAS CHARLES
TITLE OF INVENTION: MODIFIED PEPTIDES AS THERAPEUTIC AGENTS
FILE REFERENCE: A-527
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SYSTEM: PC-DOS/MS-DOS
PatentIn Release #1.0, Version #1.30
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41.5%; Score 34; DB 12;
Best Local Similarity 54.5%; Pred. No. 1.4e+02;
Matches 6; Conservative 1; Mismatches 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1; Mismatches
                                                                      CURRENT APPLICATION NUMBER: US/10/651,723
CURRENT FILING DATE: 2003-08-29
PRIOR PELLING DATE: 1999-10-22
PRIOR FILING DATE: 1999-10-22
PRIOR FILING DATE: 1998-10-23
NUMBER OF SEQ ID NOS: 1133
SEQ ID NO 783
                                                                                                                                                                                                                                                                                                                                                                                                                      ) OTHER INFORMATION: IL-1 ANTAGONIST PEPTIDE US-10-651-723-783
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: Pennie & Edmonds
STREET: 1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/602,999
FILING DATE: 16-FEB-1996
ATTORNEY/AGENT INFORMATION:
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APPLICATION NUMBER: US/10/161,791
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-10-161-791-255; Application US/10161791; Sequence 255, Application US/10161791; GENERAL INFORMATION: GENERAL INFORMATION: Applicaw B.
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                                                                                                Length 12;
                                                                                          41.5%; Score 34; DB 14; Length 12; 100.0%; Pred. No. 1.4e+02; rative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                            APPLICANT: SPARKS, Andrew B.
APPLICANT: KAY, Brian K.
APPLICANT: HORN, Judith M.
APPLICANT: QUILLIAM, Lawrence A.
APPLICANT: DER, Channing J.
APPLICANT: DER, Channing J.
APPLICANT: FUDER, James E.
TITLE OF INVENTION: SH3 BINDING PEPTIDES AND METHODS OF TITLE OF INVENTION: ISOLATING AND USING SAME
NUMBER OF SEQUENCES: 467
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: PatentIn Release #1.0, Version #1.30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: Pennie & Edmonds
STREET: 1155 Avenue of the Americas
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/602,999
FILING DATE: 16-FEB-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1101-202
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APPLICATION NUMBER: US/10/161,791
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              E: Floppy disk
IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                        ; Sequence 410, Application US/10161791; Publication No. US20030186863A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (212) 790-9090
(212) 869-9741/8864
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Misrock, S. Leslie
REGISTRATION NUMBER: 18,872
REFERENCE/DOCKET NUMBER: 110
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 410:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ouery Match
Best Local Similarity 100...
6, Conservative
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                                                                                                                                    6; Conservative
; TYPE: amino acid
; TOPDLOGY: unknown
; MOLECULE TYPE: peptide
US-10-161-791-255
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY: U.S.A.
ZIP: 10036-2711
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New York
: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: amino acid
TOPOLOGY: unknow
                                                                                            Query Match
Best Local Similarity
                                                                                                                                                                        9 PRPLPV 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                9 PRPLPV 14
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                                                                                                                                                                                                                                                                                    US-10-161-791-410
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Gaps
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SPARKS, ANDREW B.

THORN, UDDITH M.

QUILLIAM, LAWRENCE A.

DER, CHANNING J.

TITLE OF INVENTION: SLC SH3 BINDING PEPTIDES AND METHODS OF

ISOLATING AND USING SAME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
ADDRESSEE: OBLON, SPIVAK, MCCLELLAND, MAIER & NEUSTADT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   .
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                                                                                           APPLICANT: Hedley, Marzneen
APPLICANT: Hedley, Mary Lynne
APPLICANT: Tedley, Mary Lynne
APPLICANT: Tomlinson, Andrew J.
APPLICANT: Tomlinson, Andrew J.
APPLICANT: Tomlinson, Andrew J.
APPLICANT: Cole, Geoffrey
TITLE OF INVENTION: CYPIBI NUCLEIC ACIDS AND METHODS OF USE
FILE REFERENCE: 08191-021001
CURRENT APPLICATION NUMBER: US/09/999,686
CURRENT FILING DATE: 2001-10-31
PRIOR APPLICATION NUMBER: 60/298,428
PRIOR APPLICATION NUMBER: 60/298,428
PRIOR APPLICATION NUMBER: 60/298,428
PRIOR PILING DATE: 2001-01-12
PRIOR APPLICATION NUMBER: 60/244,501
PRIOR FILING DATE: 2001-01-12
PRIOR PILING DATE: 2001-01-31
PRIOR FILING DATE: 2001-01-31
SOFTWARE OF SEC FOR WINGOWS Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          P.C. STREET: 1755 S. Jefferson Davis Highway, Suite 400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       40.2%; Score 33; DB 10; Length 9; 66.7%; Pred. No. 9.7e+05; Live 1; Mismatches 2; Indels
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Villacorta, Gilberto M.
REGISTRATION NUMBER: 34,038
REFERENCE/DOCKET NUMBER: 4980-007-0
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703) 413-3000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/938,315
FILING DATE: 23-Aug-2001
CLASSIFICATION: <Unknown>
                    ; Sequence 22, Application US/09999686
; Publication No. US20030028000A1
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 86, Application US/09938315
Patent No. US20020091085A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES: 106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity 66.7
Matches 6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STATE: Virginia
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: Arlington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: PRT
CRGANISM: Homo sapiens
US-09-999-686-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    6 FVSPRPLPV 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 FLDPRPLTV 9
US-09-999-686-22
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40.2%; Score 33; DB 9; Length 13;
Best Local Similarity 62.5%; Pred. No. 2.1e+02;
Matches 5; Conservative 3; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      h 40.2%; Score 33; DB 14; Length 13; Similarity 62.5%; Pred. No. 2.1e+02; 5; Conservative 3; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: KAY, Brian K.
APPLICANT: THORN, Judith M.
APPLICANT: THORN, Judith M.
APPLICANT: OUILLIAM, Lawrence A.
APPLICANT: FOWLKES, Dana M.
APPLICANT: FOWLKES, Dana M.
APPLICANT: RIBER, James E.
TITLE OF INVENTION: SH3 BINDING PEPTIDES AND METHODS OF
NUMBER OF SEQUENCES: 467
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/10/161,791
                                                                                                                                   TOPOLOGY: unknown
MOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: SEQ ID NO: 86:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
APPLICATION NUMBER:
STILNG DATE: 16-FEB-1996
ATTORNEY/AGENT INFORMATION:
NAME: Misrock, S. Lefelie
REGISTRATION NUMBER: 18,872
REFERENCE/DOCKET NUMBER: 1101-202
TELECHONE: (212) 790-9090
TELECHANKE: (212) 790-9090
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pennie & Edmonds
1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 86, Application US/10161791
Publication No. US:20030186863A1
GENERAL INFORMATION:
APPLICANT: SPARKS, Andrew B.
TELEFAX: (703) 413-2220
TELEX: 248855 OPAT UR
INFORMATION FOR SEQ ID NO: 86:
SEQUENCE CHARACTERISTICS:
LENGTH: 13 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 86:
SEQUENCE CHARACTERISTICS:
                                                                                                                  TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 13 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                            8 SPRPLPW 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 12
US-10-161-791-86
                                                                                                                                                                                                    US-09-938-315-86
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-10-161-791-86
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Gaps
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ADDRESSEE: OBLON, SPIVAK, MCCLELLAND, MAIER & NEUSTADT,
                                                                                                                                                                                                                                                                                 QUILLIAM, LAWRENCE A.
DER, CHANNING J.
TITLE OF INVENTION: Src SH3 BINDING PEPTIDES AND METHODS OF ISOLATING AND USING SAME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  .
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET: 1755 S. Jefferson Davis Highway, Suite 400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IEBM PC compatible
COMPUTER: IEBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/938,315
FILING DATE: 23-Aug-2001
CLASSIFICATION: CURROMAN-
ATTORNEY/AGENT INFORMATION:
NAME: Villacorta, Gilberto M.
REGISTRATION NUMBER: 34,038
REFERENCE/DOCKET WUMBER: 4980-007-0
TELEPHONE: (703) 413-3200
TELEPRA: (703) 413-3220
TELEPRA: (703) 413-2220
TELERA: 248855 OPAT UR
INFORMATION FOR SEQ ID NO: 62:
SEQUENCE CHARACTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 40.2%; Score 33; DB 9; Length 14; Best Local Similarity 62.5%; Pred. No. 2.2e+02; Matches 5; Conservative 3; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY: unknown
MOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: SEQ ID NO: 62:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SPARKS, Andrew B.
KAY, Brian K.
THORN, Judith M.
OUILLIAM, Lawrence A.
DER, Channing J.
FOWLKES, Dana M.
                                                                                                                                                  Sequence 62, Application US/09938315
Patent No. US20020091085A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-10-161-791-62
Sequence 62, Application US/10161791
Publication No. US20030186863A1
GENERAL INFORMATION:
                                                                                                                                                                                                                     APPLICANT: KAY, BRIAN K.
SPARKS, ANDREW B.
THORN, JUDITH M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 14 amino acids
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                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STATE: Virginia
COUNTRY: U.S.A.
ZIP: 22202
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CITY: Arlington
SPRPLPVV 15
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                                       2 TPRPLPML 9
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APPLICANT:
APPLICANT:
APPLICANT:
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APPLICANT:
                                                                                                          RESULT 13
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Best Local Similarity

Matches

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ADDRESSEE: OBLON, SPIVAK, MCCLELLAND, MAIER & NEUSTADT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: SEC SH3 BINDING PEPTIDES AND METHODS OF ISOLATING AND USING SAME
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TITLE OF INVENTION: SH3 BINDING PEPTIDES AND METHODS OF TITLE OF INVENTION: SH3 BINDING PEPTIDES AND METHODS OF TITLE OF INVENTION: ISOLATING AND USING SAME OF SEQUENCES: 467
CORRESPONDENCE ADDRESS: ADDRESSE: Pennie & Edmonds STREET: 1155 Avenue of the Americas CITY: New York STATE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
40.2%; Score 33; DB 14; Length 14;
Best Local Similarity 62.5%; Pred. No. 2.2e+02;
Matches 5; Conservative 3; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/10/161,791
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/602,999
FILING DATE: 16-FEB-1996
ATTORNEY/AGENT INFORMATION:
NAME: Misrock, S. Leblie
REGISTRATION NUMBER: 18,872
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: KAY, BRIAN K.
PRELICANT: KAY, BRIAN K.
THORN, JUDITH M.
QUILLIAM, LAWRENCE A.
DER, CHANNING J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RETERENCE/DOCKET NUMBER: 1101-202
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 790-9090
TELEFAX: (212) 869-9741/8864
                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 67, Application US/09938315
Patent No. US20020091085A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEX: 6641 PENNIE
INFORMATION FOR SEQ ID NO: 62:
SEQUENCE CHARACTERISTICS:
LENGTH: 14 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES: 106
                                                                                                                                                                                                                                                      ZIP: 10036-z/1.
COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CITY: Arlington
STATE: Virginia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; TOPOLOGY: unknown; MOLECULE TYPE: peptide US-10-161-791-62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            8 SPRPLPVV 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               amino acid
                                                                                                                                                                                                               STATE: New York COUNTRY: U.S.A.
                                                                                                                                                                                                                                    COUNTRY: U.S.A. ZIP: 10036-2711
                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 15
US-09-938-315-67
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2, 2004, 10:11:14; Search time 32.3333 Seconds (without alignments) 121.720 Million cell updates/sec
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/ Cgn2_6/ptodata/1/pubpaa/US07_NEW_PUB.ppp:*
/ Cgn2_6/ptodata/1/pubpaa/US06_NEW_PUB.ppp:*
/ Cgn2_6/ptodata/1/pubpaa/US06_PUBCOMB.ppp:*
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/ Cgn2_6/ptodata/1/pubpaa/US08_NEW_PUB.ppp:*
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/ Cgn2_6/ptodata/1/pubpaa/US10B_PUBCOMB.ppp:*
/ Cgn2_6/ptodata/1/pubpaa/US10B_PUBCOMB.ppp:*
/ Cgn2_6/ptodata/1/pubpaa/US10B_PUBCOMB.ppp:*
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GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1070241 seqs, 262374223 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Published_Applications_AA:*
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                                                                 OM protein - protein search, using sw model
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BLOSUM62
Gapop 10.0 , Gapext 0.5
                                                                                                                                                                                                                                                                                                                                 US-10-009-049-2
79
1 SRAHSFGSPRPLPVV 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Minimum DB seq length: 0
Maximum DB seq length: 15
                                                                                                                                                                                                   April
                                                                                                                                                                                                                                                                                                                                                                    Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Database :
                                                                                                                                                                                                                                                                                                                                                                                                           Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Searched:
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

	Description	Seguence 92, Appl	Sequence 92, Appl	Sequence 17, Appl		Sequence 255, App	Sequence 410, App	Segmence 86, Appl	Sequence 86, Appl	Sequence 62, Appl	Sequence 62. Appl	Segmence 67, Appl	Sequence 67. Appl	Sequence 114. App	Sequence 4631. An	Sequence 30, Appl
	ID	US-09-938-315-92	US-10-161-791-92	US-09-945-917-17	US-09-845-917A-17	US-10-161-791-255	US-10-161-791-410	US-09-938-315-86	US-10-161-791-86	US-09-938-315-62	US-10-161-791-62	US-09-938-315-67	US-10-161-791-67	US-09-815-837-114	US-10-149-138-4631	US-10-286-457-30
	DB	6	14	10	10	14	14	δ	14	σ	14	σ	14	σ	15	14
	Query Match Length DB	13	13	Ξ	11	12	15	13	13	14	14	14	14	6 0	11	12
φo	Query Match	44.3	44.3	43.0	43.0	43.0	43.0	41.8	41.8	41.8	41.8	39.2	39.5	38.0	38.0	38.0
	Score	35	35	34	34	34	34	33	33	33	33	31	31	30	30	30
	Result No.	н	7	e	4	D.	9	7	80	თ	10	11	12	13	14	15

Sequence 30, Appl

Sequence 281, Appl Sequence 85, Appl Sequence 87, Appl Sequence 37, Appl Sequence 2561, Ap Sequence 261, Ap Sequence 311, Appl Sequence 311, Appl Sequence 311, Appl Sequence 101, Appl Sequence 111, Appl Sequence 111, Appl Sequence 111, Appl Sequence 111, Appl	144, 181, 181, 181, 181, 181, 181, 181,	4440226	
0110-100-100-100-100-100-100-100-100-10	09-572-404B-13 09-572-404B-13 10-044-896-9 09-291-417-78 09-291-417-114 10-153-756-14	10-15/-7/5B 10-254-446A 10-155-477- 10-185-050- 10-161-791- 10-008-524A	US-10-008-524A-27 US-10-008-524A-28 US-10-008-524A-29 US-10-350-719-26 US-10-350-719-27
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ALIGNMENTS

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QUILLIAM, LAWRENCE A.
DER, CHANNING J.
TITLE OF INVENTION: Src SH3 BINDING PEPTIDES AND METHODS OF ISOLATING AND USING SAME
                                                                                                                                                                                                                                                                                                                                             ADDRESSEE: OBLON, SPIVAK, MCCLELLAND, MAIER & NEUSTADT,
                                                                                                                                                                                                                                                                                                                                                                                        STREET: 1755 S. Jefferson Davis Highway, Suite 400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/938,315
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION: CONCURS CLASSIFICATION: THORMATION:
NAME: Villacorta, Gilberto M.
REGISTRATION NUMBER: 34,038
REPERENCE/DOCKET NUMBER: 4980-007-0
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703) 413-3000
                                             ; Sequence 92, Application US/09938315; Patent No. US20020091085A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: 23-Aug-2001
                                                                                                                  KAY, BRIAN K.
SPARKS, ANDREW B.
THORN, JUDITH M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: (703) 413-2220
TELEX: 248855 OPAT UR
                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES: 106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ZIP: 22202
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                       CITY: Arlington
                                                                                                                                                                                                                                                                                                                                                                                                                                           STATE: Virginia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY: U.S.A.
                                                                                                                       APPLICANT: KAY
RESULT 1
US-09-938-315-92
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                                                                                                                                                                                                        Query Match

44.3%; Score 35, DB 9, Length 13;
Best Local Similarity 62.5%; Pred. No. 1.1e+02;
Matches 5; Conservative 3; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: SH3 BINDING PEPTIDES AND METHODS OF TITLE OF INVENTION: ISOLATING AND USING SAME NUMBER OF SEQUENCES: 467
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ZIP: 10036-2711

ZIP: 10036-2711

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                             TYPE: amino acid
TOPOLOGY: unknown
MOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: SEQ ID NO: 92:
US-09-938-315-92
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: Pennie & Edmonds
STREET: 1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/602,999
FILING DATE: 16-FEB-1996
ATTORNBY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Misrock, S. Lealie
REGISTRATION NUMBER: 18,872
REFERENCE/DOCKET NUMBER: 1101-202
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/10/161,791
                                                                                                                                                                                                                                                                                                                                                                                                                                  US-10-161-791-92; Sequence 92, Application US/10161791; Publication No. US20030186863A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: QUILLIAM, Lawrence A. APPLICANT: DER, Channing J. APPLICANT: POWIKES, Dana M. APPLICANT: RIDER, James E.
INFORMATION FOR SEQ ID NO: 92:
SEQUENCE CHARACTERISTICS:
LENGTH: 13 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEPHONE: (212) 790-9090
TELEFAX: (212) 869-9741/8864
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 92:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Andrew B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: SPARKS, Andrew B
APPLICANT: KAY, Brian K.
APPLICANT: THORN, JUGITH M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 62.5
Matches 5; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; MOLECULE TYPE: peptide US-10-161-791-92
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                 8 SPRPLPWV 15
                                                                                                                                                                                                                                                                                                                                           2 TPRPLPLI 9
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: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE:
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY:
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Publication No. US20030167538A1
GENERAL INFORMATION:
APPLICAMY: BOGAER, Thierry
APPLICAMY: BOGAER, Thierry
APPLICAMY: Wandekerckhove, Joel
TITLE OF INVENTION: PRACESSES FOR THE IDENTIFICATION OF COMPOUNDS WHICH
TITLE OF INVENTION: OWTROL CELL BEHAVIOUR, THE COMPOUNDS IDENTIFIED AND
TITLE OF INVENTION: USE IN THE CONTROL OF CELL BEHAVIOUR
FILE REFERENCE: P/14-1
CURRENT APPLICATION NUMBER: US/09/845,917A
CURRENT PLING DATE: 201-04-30
NUMBER OF SEQ ID NOS: 58
SOFTWARE: Patentin Ver. 2.1
                                                                                                                APPLICANT: BOGGER, Thierry
APPLICANT: BOGGER, Thierry
APPLICANT: Bogger, Joel
TITLE OF INVENTION: PROCESSES FOR THE IDENTIFICATION OF COMPOUNDS WHICH
TITLE OF INVENTION: CONTROL CELL BEHAVIOUR, THE COMPOUNDS IDENTIFIED AND
TITLE OF INVENTION: PHARMACEUTICAL COMPOSITIONS CONTINUING THEM AND THEIR
FILE OF INVENTION: USE IN THE CONTROL OF CELL BEHAVIOUR
FILE REFERENCE: P/14-1
CURRENT APPLICATION NUMBER: US/09/945,917
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 11;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
43.0%; Score 34; DB 10; L
Best Local Similarity 100.0%; Pred. No. 1.4e+02;
Matches 6; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 255, Application US/10161791; Publication No. US20030186863A1
GENERAL INFORMATION: APPLICANT: SPREKS, Andrew B. APPLICANT: KAY, Brian K.
                                                ; Sequence 17, Application US/09945917; Publication No. US20030042381A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ) ORGANISM: Caenorhabditis elegans US-09-945-917-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: PRT ORGANISM: Caenorhabditis elegans
                                                                                                                                                                                                                                                                                                                      CURRENT FILING DATE: 1998-09-21
NUMBER OF SEQ ID NOS: 58
SOFTWARE: PatentIn Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    6; Conservative
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Best Local Similarity
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                                                                                              GENERAL INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 4
US-09-845-917A-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-09-845-917A-17
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RESULT 3
US-09-945-917-17
                                                                                                                                                                                                                                                                                                                                                                                              SEQ ID NO 17
LENGIH: 11
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LENGTH: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 5
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Gaps 0;

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Publication No. U520030186863A1

GENERAL INFORMATION:
APPLICANT: SPARKS, Andrew B.
APPLICANT: THORN, Judith M.
APPLICANT: QUILLIAM, Lawrence A.
APPLICANT: QUILLIAM, Lawrence A.
APPLICANT: POWIKES, Dana M.
APPLICANT: POWIKES, Dana M.
APPLICANT: STERR, James E.
TITLE OF INVENTION: SH3 BINDING PEPTIDES AND METHODS OF TITLE OF INVENTION: SH3 BINDING SAME
MINDER OF SECTION SHADES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       43.0%; Score 34; DB 14; Length 12; 100.0%; Pred. No. 1.5e+02; tive 0; Mismatches 0; Indels
                  APPLICANT: QUILLIAM, Lawrence A.
APPLICANT: DER, Channing J.
APPLICANT: FOWLKES, Dana M.
APPLICANT: FIDER, James E.
TITLE OF INVENTION: SH3 BINDING PEPTIDES AND METHODS OF
TITLE OF INVENTION: ISOLATING AND USING SAME
NUMBER OF SEQUENCES: 467
                                                                                                                                                                                                                                                                                        COUNTRY: U.S.A.
ZIP: 10056-2711
COMPUTER READABLE PORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/161,791
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES: 467
CORRESPONDENCE ADDRESS:
CADDRESSES: Pennie & Edmonds
STREET: 1155 Avenue of the Americas
                                                                                                                                                                               CORRESPONDENCE ADDRESS:
ADDRESSEE: Pennie & Edmonds
STREET: 1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTOKNE 1, ANGEL.

NAME: Mistock, S. Leslie
REGISTRATION NUMBER: 18,872
REFERENCE/DOCKET NUMBER: 1101-202
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 869-9741/8864
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/602,999
FILING DATE: 16-FEB-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 255:
SEQUENCE CHARACTERISTICS:
THORN, Judith M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity 100 (
Matches 6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 12 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: unknown
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          amino acid
                                                                                                                                                                                                                                                                            New York
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                                                                                                                                                                                                                                                  New York
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US-10-161-791-410
  APPLICANT:
                                                                                                                                                                                                                                                    CITY: 1
STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STATE:
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Gaps
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SPARKS, ANDREW B.
THORN, JUDITH M.
QUILLIAM, LIAMRENCE A.
DER, CHANNING J.
TITLE OF INVENTION: STC SH3 BINDING PEPTIDES AND METHODS OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES: 106
CORRESPONDENCE ADDRESS:
ADDRESSEE: OBLON, SPIVAK, MCCLELLAND, MAIER & NEUSTADT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            .
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STREET: 1755 S. Jefferson Davis Highway, Suite 400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          43.0%; Score 34; DB 14; Length 15; 100.0%; Pred. No. 1.8e+02; Live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC_formatible
OPERATING SYSTEM: PC_DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                        SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/161,791
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/938,315
FILING DATE: 23-Aug-2001
CLASSIFICATION: <UNKnown.
ATTORNEY/AGENT INFORMATION:
NAME: VIllacorta, Gilberto M.
REGISTRATION NUMBER: 34,038
REFERENCE/DOCKET NUMBER: 4980-007-0
                                                                                                                                                   FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/602,999
FILING DATE: 16-FEB-1996
ATTORNEY/AGENT INFORMATION:
NAME: Misrock, S. Leslie
REGISTRATION NUMBER: 18,872
REFERENCE/DOCKET NUMBER: 1101-202
TELECOMMUNICATION INFORMATION:
                                             COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 86, Application US/09938315
Patent No. US20020091085A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                   TELEPHONE: (212) 790-9090
TELEFAX: (212) 869-9741/8864
TELEX: 66141 PENNER
INFORMATION FOR SEQ ID NO: 410:
SEQUENCE CHARACTERISTICS:
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 100...
Fra 6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 15 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CITY: Arlington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STATE: Virginia COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; MOLECULE TYPE: peptide US-10-161-791-410
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: amino acid
TOPOLOGY: unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     9 PRPLPV 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                4 PRPLPV 9
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Gaps

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3; Mismatches

5; Conservative

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Matches
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                                                                                                                                                                                                                                                          Length 13;
                                                                                                                                                                                                                                                                                                   0: Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: THON, Judith M.
APPLICANT: THON, Judith M.
APPLICANT: OUILLIAM, Lawrence A.
APPLICANT: FOWLKES, Dana M.
APPLICANT: FUBER, James E.
APPLICANT: RIDER, James E.
TITLE OF INVENTION: SH3 BINDING PEPTIDES AND METHODS OF TITLE OF INVENTION: ISOLATING AND USING SAME
NUMBER OF SEQUENCES: 467
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARR: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                      Query Match 41.8%; Score 33; DB 9; I
Best Local Similarity 62.5%; Pred. No. 2.2e+02;
Matches 5; Conservative 3; Mismatches 0,
                                                                                                                                              TOPOLOGY: unknown
MOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: SEQ ID NO: 86:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
APPLICATION NUMBER:
FILING DATE: 16 FEB-1996
ATTORNEY/AGENT INFORMATION:
NAME: Misrock, S. Leslie
REGISTATION NUMBER: 18,872
REFERENCE/DOCKET NUMBER: 1101-202
TELECOMMUNICATION INPORMATION:
TELECOMMUNICATION INPORMATION:
TELECHNOME: (212) 790-9090
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    E: Pennie & Edmonds
1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/10/161,791
TELEPHONE: (703) 413-3000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 86, Application US/10161791
Publication No. US20030186863A1
GENERAL INFORMATION:
               TELEFAX: (703) 413-2220
TELEX: 248855 OPAT UR
                                                 INFORMATION FOR SEQ ID NO: 86:
SEQUENCE CHARACTERISTICS:
LENGTH: 13 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (212) 869-9741/8864
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 86:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SPARKS, Andrew B.
                                                                                                                         TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 13 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    KAY, Brian K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY: U.S.A.
ZIP: 10036-2711
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                         8 SPRPLPVV 15
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2 TPRPLPML 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 8
US-10-161-791-86
                                                                                                                                                                                                               US-09-938-315-86
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-10-161-791-86
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGIH:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
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Best Local Similarity

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Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENČE ADDRESS:
ADDRESSEE: OBLON, SPIVAK, MCCLELLAND, MAIER & NEUSTADT,
                                                                                                                                                                                                                                                                                                       QUILLIAM, LAWRENCE A.

DER, CHANNING J.

TITLE OF INVENTION: Src SH3 BINDING PEPTIDES AND METHODS OF ISOLATING AND USING SAME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   .
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
41.8%; Score 33; DB 9; Length 14;
Best Local Similarity 62.5%; Pred. No. 2.4e+02;
Matches 5; Conservative 3; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/938,315
FILING DATE: 23-Aug-2001
CLASSIFICATION: <UDAKnown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
NAME: Villacorta, Gilberto M.
REGISTRATION NUMBER: 34,038
REFERENCE/DOCKET NUMBER: 4980-007-0
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703) 413-3000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MG-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: unknown

MOLECULE TYPE: peptide

SEQUENCE DESCRIPTION: SEQ ID NO: 62:
US-09-938-315-62
                                                                                                                                                              Sequence 62, Application US/09938315
Patent No. US20020091085A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; Sequence 62, Application US/10161791
; Publication No. US20030186863A1
; GENBRAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: SPARKS, Andrew B.
APPLICANT: KAY, Brian K.
APPLICANT: THORN, Judith M.
APPLICANT: QUILLIAM, Lawrence A.
APPLICANT: DER, Channing J.
                                                                                                                                                                                                                                                          SPARKS, ANDREW B.
THORN, JUDITH M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: (703) 413-2220
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 14 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEX: 248855 OPAT UR
INFORMATION FOR SEQ ID NO: 62:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                    BRIAN K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CITY: Arlington
STATE: Virginia
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES:
8 SPRPLPW 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             8 SPRPLPVV 15
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                             :|||||:
2 TPRPLPML 9
                                                                                                                                                                                                                                      APPLICANT: KAY,
                                                                                                                                             US-09-938-315-62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-10-161-791-62
                                                                                                                    RESULT 9
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us-10-009-049-2.szlm15.rapb

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TELEPHONE: (212) 790-9090
TELEFAX: (212) 869-9741/8864
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                5; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sest Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        8 SPRPLP 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    :|||||
3 TPRPLP 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STATE: New York
COUNTRY: U.S.A.
ZIP: 10036-2711
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 12
US-10-161-791-67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AL INFORMATION:
APPLICANT: KAY, BRIAN K.
SPARKS, ANDREW B.
THORN, JUDITH M.
QUILLIAM, LAWRENCE A.
DER, CHANNING J.
TITLE OF INVENTION: SIC SH3 BINDING PEPTIDES AND METHODS OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
ADDRESSEE: OBLON, SPIVAK, MCCLELLAND, MAIER & NEUSTADT,
P.C.
STREET: 1755 S. Jefferson Davis Highway, Suite 400
CITY: Arlington
STATE: Virginia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            .
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                41.8%; Score 33; DB 14; Length 14; 62.5%; Pred. No. 2.4e+02; ive 3; Mismatches 0; Indels
APPLICANT: FOWLKES, Dana M.
APPLICANT: RIDER, James B.
TITLE OF INVENTION: SH3 BINDING PEPTIDES AND METHODS OF
TITLE OF INVENTION: ISOLATING AND USING SAME
NUMBER OF SUCHENCES: 467
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                             MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/161,791
                                                                                                                     RESSEE: Pennie & Edmonds
ET: 1155 Avenue of the Americas
E: New York
E: New York
TRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION UNDRER: US/08/602,999
PILING DATE: 16-FEB-1996
ATTORNEY/AGENT INFORMATION:
NAME: Misrock, S. Leslie
REGISTRATION NUMBER: 18,872
REFERENCE/DOCKET NUMBER: 1101-202
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 67, Application US/09938315
Patent No. US20020091085A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEPHONE: (212) 790-9090
TELEFAX: (212) 869-9741/8864
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 62:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 14 amino acids
TYPE: amino acid
TOPOLOGY: unknown
MOLECULE TYPE: peptide
US-10-161-791-62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 62.5
Matches 5; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY: U.S.A.
                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             8 SPRPLPWV 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3 TPRPLPML 10
                                                                                                                       ADDRESSEE:
STREET: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-09-938-315-67
                                                                                                                                                                                                       COUNTRY:
                                                                                                                                                               CITY: 1
STATE:
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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: KAY, Brian K.
APPLICANT: THORN, Judith M.
APPLICANT: THORN, Judith M.
APPLICANT: QUILLIAM, Lawrence A.
APPLICANT: DER, Channing J.
APPLICANT: FOWLKES, Dana M.
APPLICANT: RIBER, James B.
TITLE OF INVENTION: SH3 BINDING PEPTIDES AND METHODS OF TITLE OF INVENTION: ISOLATING AND USING SAME
NUMBER OF SEQUENCES: 467
CORRESPONDENCE ADDRESS:
             SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/938,315
FILING DATE: 23-Aug-2001
CLASSIFICATION: <UNKNOWN>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/161,791
                                                                                                                                            ATTORNEY AGENT INFORMATION:
NAME: Villacorta, Gilberto M.
REGISTRATION NUMBER: 34,038
REENCE/DOCKET NUMBER: 4980-007-0
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703) 413-3000
TELEX: 248855 OPPT UR
INFORMATION FOR SEQ ID NO: 67:
SEQUENCE CHARACTERISTICS:
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                           ; TOPOLOGY: unknown; MOLECULE TYPE: peptide; SEQUENCE DESCRIPTION: SEQ ID NO: 67: US-09-938-315-67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRICR APPLICATION DATA:
APPLICATION NUMBER: US/08/602,999
FILING DATE: 16-FEB-1996
ATTORNEY AGENT INFORMATION:
NAME: Migrock, S. Leglie
REGISTATION NUMBER: 18,872
REFERENCE/DOCKET NUMBER: 1101-202
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET: 1155 Avenue of the Americas CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 67, Application US/10161791
Publication No. US2030186863A1
GENERAL INFORMATION:
APPLICANT: SPARKS, Andrew B.
                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 14 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE: Pennie & Edmonds
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APPLICANT: Celis, Esteban
APPLICANT: Keogh, Elissa
APPLICANT: Keogh, Elissa
TITLE OF INVENTION: Inducing Cellular Immune Responses to
TITLE OF INVENTION: HER2/neu Using Peptide and Nucleic Acid Compositions
FILE REFERENCE: 2060.0140001
                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FRATURE: OTHER INFORMATION: Description of Artificial Sequence:downstream OTHER INFORMATION: linker for COS85 and COS92
                                                                                                                                                                                                               39.2%; Score 31; DB 14; Length 14; 83.3%; Pred. No. 4.8e+02; tive 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Zhu, Shirley
APPLICANT: Zhu, Shirley
APPLICANT: Arimilli, Subhashini
APPLICANT: Wang, Aljun
APPLICANT: Wang, Aljun
APPLICANT: Corixa Corporation
ITILE OF INVENTION: Immune Medators and Related Methods
FILE REFERENCE: 014058-005670US
CURRENT APPLICATION NUMBER: US/20/915,837
CURRENT FILING DATE: 2001-03-22
FRIOR APPLICATION NUMBER: US 60/191,274
FRIOR APPLICATION NUMBER: US 60/204,249
FRIOR FILING DATE: 2000-05-15
FRIOR FILING DATE: 2000-05-15
FRIOR FILING DATE: 2001-01-23
NUMBER OF SEQ ID NOS: 129
SOFTWARE: PatentIn Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity 62.5%; Pred. No. 9.7e+05; Similarity 62.5%; Pred. No. 9.7e+05; 5. Conservative 1; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-10-149-138-4631

Sequence 4631, Application US/10149138

Publication No. US20040018971A1

GENERAL INFORMATION:

APPLICANT: Fikes, John

APPLICANT: Setter, Aleasandro

APPLICANT: Southwood, Scott
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 114, Application US/09815837
Patent No. US20020082411A1
                        67:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: PRT
ORGANISM: Artificial Seguence
                                                                     14 amino acida
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                   Best Local Similarity 83.3
Matches 5; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Carter, Darrick
                                                                                                                TOPOLOGY: unknown
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           6 FGSPRPLP 13
                                                                                            amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity
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1 FDAPSPLP 8
                                                                                                                                                                                                                                                                                                              8 SPRPLP 13
                                                                                                                                                                                                                                                                                                                                         :|||||
3 TPRPLP 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-09-815-837-114
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                                                                                                                                                                  US-10-161-791-67
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LENGTH: 8
                                                                        LENGTH:
                                                                                                                                                                                                                 Query Match
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                                                                                              TYPE:
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OTHER INFORMATION: artificial sequence isolated from random peptide libraries, based OTHER INFORMATION: ability to selectively bind to endothelial cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 30, Application US/10286457

Sequence 30, Application US/10286457

GENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INVENTION: RENOTHELIAL-CELL BINDING PEPTIDES FOR DIAGNOSIS AND THERAPY
FILE REPERENCE: GFOI-P01-178
CURRENT APPLICATION NUMBER: US/10/286,457

CURRENT FILING DATE: 2002-11-01

PRIOR PILING DATE: 2001-11-01

NUMBER OF SEQ ID NOS: 684
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                                                                                                                                                                                                                                                                                                                                                        Length 11;
                                                                                                                                                                                                                                                                                                                                                                                               1; Indels
                                                                                                                                                                                                                                                                                                                                                    38.0%; Score 30; DB 15;
63.6%; Pred. No. 5.3e+02;
tive 1; Mismatches 1,
PRIOR APPLICATION NUMBER: 2002-06-10
PRIOR APPLICATION NUMBER: PCT/USO0/33591
PRIOR FILING DATE: 2000-12-11
PRIOR APPLICATION NUMBER: US 09/458,299
PRIOR FILING DATE: 1999-12-11
NUMBER OF SEQ ID NOS: 4641
SEQ ID NO 4631
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2, 2004, 10:23:35
                                                                                                                                                                                                                                                                                   ; OTHER INFORMATION: Artificial Peptide US-10-149-138-4631
                                                                                                                                                                                                                                          ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: Patentin version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 60.0°
Matches 6; Conservative
                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity 63.6
Matches 7; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                         6 FGSPR--PLPV 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 FPSPREGPLPI 11
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US-10-286-457-30
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LENGTH: 12
                                                                                                                                                                                                                     TYPE: PRT
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GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
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OM protein - protein search, using sw model

April Run on:

2, 2004, 09:56:18 ; Search time 46 Seconds (without alignments) 92.135 Million cell updates/sec

US-10-009-049-3

1 CRAHSFVSPRPLPVV 15 Perfect score: Sequence: BLOSUM62 Gapop 10.0 , Gapext 0.5 Scoring table:

1586107 seqs, 282547505 residues Searched:

447529 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 15

Listing first 45 summaries Maximum Match 100% Post-processing: Minimum Match 0%

geneseqp2000s:* geneseqp2001s:* geneseqp2002s:* geneseqp2003as:* geneseqp2003bs:* geneseqp2004s:* A_Geneseq_29Jan04:* geneseqp1980s;* geneseqp1990s;* Database:

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

	Description	Aab59251 Peptide #	6	0	9 Proline-	'n		Aam12950 HLA-A26 d	Aam08833 HLA-A1 de	9 HLA-B *	1 Src Si	Aaw05098 Proline-r	Aaw94619 Src homol	Aam09403 HLA-Al de	Aam13226 HLA-A26	m	_	Aam09490 HLA-B *07	Aaw16127 Peptide o	Aaw68767 Peptide h	Aaw58254 Interleuk	Aay09898 Interleuk	6	Abb72575 Interleuk	Aaw39011 Peptide 1	Aam11755 HLA-B *15
SUMMARIES	ID	AAB59251	AAB59249	AAB59250	AAW05069	AAM06985	AAM08985	AAM12950	AAM08833	AAM09489	AAW11111	AAW05098	AAW94619	AAM09403	AAM13226	AAM11873	AAM09177	AAM09490	AAW16127	AAW68767	AAW58254	AAY09898	AAB17679	ABB72575	AAW39011	AAM11755
	DB	4	4	4	7	4	4	4	4	4	7	N	ď	4	4	4	4	4	~	7	~	~	3	5	7	4
	Length	15	15	15	15	10	10	70	70	10	13	13	6	10	10	10	10	10	12	12	12	12	12	12	15	6
₩	Query	100.0	91.5	80.5	47.6	45.1	45.1	45.1	45.1	45.1	42.7	42.7	41.5	•	41.5	•	41.5	•	٠	41.5	41.5	41.5	41.5	41.5	41.5	40.2
	Score			99			37	37	37		35	35	34	34	34	34	34	34	34	34	34	34	34	34	34	33
	Result No.	Н	7	æ	4	5	9	7	80	თ	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25

-	Aam07008 HLA-A *02	Aam06907 Cytochrom	Aam08918 HLA-B *27	Aam06958 CYP1B1 HL	Aam11125 HLA-B8 no	Aam11262 HLA-A26 n	Aam07025 HLA-Al no	Aam06929 HLA-A2 pr	Abg68876 Human CYP	-	Aar93450 GST-PI3K	Aar93395 GST-FYN S	Aaw11098 Src SH3 d	Aaw25512 SH3 synth	Aam53295 Human non	Aap60604 Sequence	Aam09704 HLA-B8 oc	Aam08460 HLA-B *27	Aam12347 HLA-A *02
AAM07477	AAM07008	AAM06907	AAM08918	AAM06958	AAM11125	AAM11262	AAM07025	AAM06929	ABG68876	AAW11105	AAR93450	AAR93395	AAW11098	AAW25512	AAM53295	AAP60604	AAM09704	AAM08460	AAM12347
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σ	σ	σ	o,	σ	0	σ	0	σ	0	13	14	14	14	14	14	14	œ	σ	σ
40.2	40.2	40.2	40.2	40.2	40.2	40.2	40.2	40.2	40.2	40.2	40.2	40.2	40.2	40.2	40.2	39.0	37.8	37.8	37.8
33	33	33	33	33	33	33	33	33	33	33	33	33	33	33	33	32	31	31	31
26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45

ALIGNMENTS

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Fibrin fragment E; cell proliferation; cancer; vascular restenosis; arthritis; diabetes; renal disease; psoriasis.
                                                    Peptide #3 recognised by gene VIII library clone 45.
                                                                                                                                                                         Stirk CM;
             AABS9251 standard; peptide; 15 AA.
                                                                                                                                07-JUN-2000; 2000WO-GB002197.
                                                                                                                                              99GB-00012994.
                                       (first entry)
                                                                                                                                                                        Тһомрвол WD,
                                                                                                                                                          (UYAB-) UNIV ABERDEEN.
                                                                                                                                                                                     WPI; 2001-080581/09.
                                                                                                    WO200075175-A1.
                                                                                                                                             07-JUN-1999;
                                                                                                                   14-DEC-2000.
                                        27-MAR-2001
                                                                                                                                                                        Melvin WT,
                                                                                       Synthetic.
                          AAB59251;
RESULT 1
     AAB59251
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New peptides capable of modulating a fibrin fragment E activity, useful for inhibiting uncontrolled cell proliferation, e.g. for treating cancer, vascular restenosis, atherogenesis or psoriasis, and for wound healing.

Claim 1; Page 49; 59pp; English.

The present invention relates to a peptide or its fragment, which is capable of modulating a fibrin fragment E activity. The peptides and modulators are useful for inhibiting stimulation of cell proliferation induced by a fibrin degradation product. The antibody or binding fragment is also useful for identifying the active site of the fibrin fragment E receptor. The peptides are particularly useful in treating cancer, vascular restenosis, rheumatoid arthritis, diabetes, renal diseases or psoriasis. The antibodies are also useful for modulating the activity of the peptide

Sequence 15 AA;

Query Match

DB 4; Length 15; 100.0%; Score 82;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention relates to a peptide or its fragment, which is capable of modulating a fibrin fragment B activity. The peptides and modulators are useful for inhibiting stimulation of cell proliferation induced by a fibrin degradation product. The antibody or binding fragment is also useful for identifying the active site of the fibrin fragment ecceptor. The peptides are particularly useful in treating cancer, vascular restenosis, rheumatoid arthritis, diabetes, renal diseases or psoriasis. The antibodies are also useful for modulating the activity of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New peptides capable of modulating a fibrin fragment E activity, useful for inhibiting uncontrolled cell proliferation, e.g. for treating cancer, vascular restenosis, atherogenesis or psoriasis, and for wound healing.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                    Gaps
                                                                                                                                                                                                                                                                                             Fibrin fragment E; cell proliferation; cancer; vascular restenosis;
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                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Peptide #2 recognised by gene VIII library clone 45.
Pred. No. 1.3e-06;
Mismatches 0;
                                                                                                                                                                                                                                                            Peptide recognised by gene VIII library clone 45.
                                                                                                                                                                                                                                                                                                          arthritis, diabetes; renal disease; psoriasis.
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                                                                                                                                                               AAB59249 standard; peptide; 15 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAB59250 standard; peptide; 15 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 1; Page 49; 59pp; English
100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                         07-JUN-2000; 2000WO-GB002197.
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                15; Conservative
                                             1 CRAHSFVSPRPLPVV
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (UYAB-) UNIV ABERDEEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2001-080581/09
Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 15 AA;
                                                                                                                                                                                                                                                                                                                                                                            WO200075175-A1.
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                                                                                                                                                                                                                          27-MAR-2001
                                                                                                                                                                                              AAB59249;
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   Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                             RESULT 2
AAB59249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 3
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The present invention relates to a peptide or its fragment, which is capable of modulating a fibrin fragment E activity. The peptides and modulators are useful for inhibiting stimulation of cell proliferation induced by a fibrin degradation product. The antibody or binding fragment is also useful for identifying the active site of the fibrin fragment erceptor. The peptides are particularly useful in treating cancer, vascular restenosis, rheumatoid architis, dabetes, renal diseases or psoriasis. The antibodies are also useful for modulating the activity of
                                                                                                                                                                                                                                                                        New peptides capable of modulating a fibrin fragment B activity, useful for inhibiting uncontrolled cell proliferation, e.g. for treating cancer, vascular restenosis, atherogenesis or psoriasis, and for wound healing.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
           Fibrin fragment E; cell proliferation; cancer; vascular restenosis; arthritis; diabetes; renal disease; psoriasis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   7. .13
/label= SH3_binding_motif
/note= "corresponds to consensus sequence RPLPXXP"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Proline-rich SH3 binding peptide fuse5B-4.51 from random library.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 66; DB 4; Length 15;
Pred. No. 0.00057;
0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      domain; SH3 domain; oncogene; consensus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
                                                                                                                                                                                                                   Stirk CM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAW05069 standard; peptide; 15 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0;
                                                                                                                                                                                                                                                                                                                                 Claim 1; Page 49; 59pp; English.
                                                                                                                                    07-JUN-2000; 2000WO-GB002197.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2 RAHSFVSPRPLPVV 15
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                                                                                                                                                                                                                     Thompson WD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     random display library.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (revised)
                                                                                                                                                                                         (UYAB-) UNIV ABERDEEN
                                                                                                                                                                                                                                             WPI; 2001-080581/09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 15 AA;
                                                                               WO200075175-A1,
                                                                                                                                                               07-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   19-APR-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    13;
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                                                                                                         14-DEC-2000
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                                                                                                                                                                                                                  Melvin WT,
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(RHON ) RHONE POULENC RORER PHARM INC.
94US-00230047.
                                                                     Ricca GA,
                                                                   Ivashchenko YD, Ricca G
                                                                                                                        WPI; 1996-361965/36.
                                                                                        French SM,
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Jaye MC;

Cheadle C, South VJ,

comparison of the positive peptides revealed a highly reiterated, proline "rich," a mino acid consensus motif (i.e. the sequence RPLPXXP). This motif demonstrated a strong similarity to a Src SH3 binding sequence identified within a protein isolated by screening a lambda-lox mouse mebryo cDNA library with a Src SH3 domain probe. The present sequence is that of one of the positive peptides dientified from the fuse SB random display library. (Updated on SS-WAR-2003 to correct or field.) A fusion phage library for expressing random 15mer peptides was prepared using the filamentous bacteriophage fuse 5B. The display library was screened for peptides which bind to the Src SH3 domain. A sequence New DNA encoding proteins that bind to Src-homology domain 3 - used to inhibit signal transduction, e.g. in cases of T cell activation. Example 1; Col 7; 39pp; English.

Sequence 15 AA;

Gaps .. 0 47.6%; Score 39; DB 2; Length 15; 70.0%; Pred. No. 16; 3; Indels tive 0; Mismatches 3; Indels 7; Conservative Query Match Best Local Similarity Matches

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4 HSFVSPRPLP 13 HSHFHPRPLP 10

RESULT 5

AAM06985 standard; peptide; 10 AA. 09-OCT-2001 AAM06985; PARTY SERVER SER

CYP1B1 HLA-A2 binding peptide fragment #35.

(first entry)

Cytochrome P450 1B1; CYPlB1; cancer; immunotherapy; diagnosis; HLA; human leukocyte antigen, oyrotoxic T-lymphocyte; CTL; MHC; APC; mador histocompatibility complex; antigen presenting call; melanoma; lymphoma; carcinoma; maltiple myeloma; leukaemia.

WO200135810-A2.

25-MAY-2001

15-NOV-2000; 2000WO-US031513.

99US-0165590P. 15-NOV-1999; FARBER CANCER INST INC

UNIV BOSTON.

(UYBO-)

DAND DANA

Nadler LM, Maecker B; Sherr D, Schultze JL, Vonderheide RH, Von Bergwelt- Baildon M;

WPI; 2001-355537/37.

Treating a cancer patient involves administering to patient a cytotoxic T -lymphocyte, an antigen presenting cell that activates T cells, a peptide of cytochrome P450 1B1, or a polynucleotide encoding the peptide.

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The present any order of the present any expectation of the present any expension or is at risk of having, a cell enteresting cytochrome P450 IBI (CYPIBI), which involves administering to the patient a cytochroxic T-lymphocyte of CTL, a peptide of CYPIBI that binds to major histocompatibility activates CTL, a peptide of CYPIBI that binds to major histocompatibility complex (MHC) molecule, or a nucleic acid molecule encoding CYPIBI or complex (MHC) molecule, or a nucleic acid molecule encoding CYPIBI or complex (MHC) molecule, involving measuring the level of CTL binds to MHC complex molecule, involving measuring the level of CTL complex molecule, involving measuring the level of CTL complex molecule, involving measuring the level of CTL complex molecule. (I) that binds to MHC complex molecule, (3) an expirate of a CYPIBI peptide (1) that binds to MHC complex molecule, (3) an expirate of a CYPIBI peptide (1) that binds to MHC complex molecule. The method is useful for treating a patient or generated APC (III) that presents a peptide of a CYPIBI in the context of a MHC complex molecule. The method is useful for treating a patient of a surface or is useful for the prevention, treatment and diagnosis of cancer, e.g. complex molecule, and useful for the prevention, treatment and diagnosis of cancer, e.g. complex ovarian, uterine, cervical, prostate, liver, colon, pancreatic and brain cancer. AAM06905 to AAM1356s represent CYPIBI peptides which can bind to human leukocyte antigens (HinAs), as well as other amino acid context of the present invention acid in the exemplification of the present invention
                                                                                             The present invention describes a method for treating a patient having
                                Disclosure; Page 40; 121pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 10 AA;
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0; Gaps Score 37; DB 4; Length 10; Pred. No. 23; 1; Mismatches 2; Indels 45.18; 70.08; 7; Conservative Query Match Best Local Similarity Matches

.. 0

6 FVSPRPLPVV 15 1 FLDPRPLTVV à 셤

AAM08985 standard; peptide; 10 AA. AAM08985; RESULT 6

(first entry) 09-OCT-2001

HLA-A *0201 decamer #2.

Cytochrome P450 1B1; CYP1B1; cancer; immunotherapy; diagnosis; HLA; human leukcoyte antigen; cytochroxic T-1ymphocyte; CTL; MHC; APC; major histocompatibility complex; antigen presenting cell; melanoma; lymphoma; carcinoma; sarcoma; multiple myeloma; leukaemia.

Homo sapiens.

WO200135810-A2.

25-MAY-2001

15-NOV-2000; 2000WO-US031513

15-NOV-1999;

FARBER CANCER INST INC (DAND) DANA FARBER (UYBO-) UNIV BOSTON.

Maecker B; Vonderheide RH, Sherr D, Nadler LM, Von Bergwelt- Baildon M; Schultze JL, PART S X B X S B X

WPI; 2001-355537/37.

Treating a cancer patient involves administering to patient a cytotoxic T -lymphocyte, an antigen presenting cell that activates T cells, a peptide of cytochrome P450 1B1, or a polynucleotide encoding the peptide.

Fri Apr

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The present invention describes a method for treating a patient having, or is at risk of having, a cell expressing cytochrome P450 1B1 (CYP1B1), white which involves administering to the patient a cytocoxic T-lymphocyte (CTL) that kills the cell, an anticlen presenting cell (APC) that activates CTL, a peptide of CYP1B1 that binds to major histocompatibility complex (MHC) molecule, or a nucleic acid molecule encoding CYP1B1 or peptide of CYP1B1. Also described are: (1) a method for assessing the clevel of immunity of a patient to CYP1B1 or a peptide of CYP1B1 that binds to MHC complex molecule, involving measuring the level of CTL specific for CYP1B1 or the peptide of CYP1B1 in a sample from the patient (2) a CYP1B1 peptide (1) that binds to MHC complex molecule, (3) an exvivo generated APC (III) that specifically kills a cell expressing CYP1B1 in a specific of a MHC complex-restricted fashion; and (4) and exvince context of a MHC complex molecule. The method is useful for treating a patient context of a mHC complex molecule. The method is useful for treating a patient context of a mHC complex molecule. The method is useful for treating a patient context of a mHC complex molecule. The method is useful for treating a patient context of a mHC complex molecule. The method is useful for treating a patient context of a min overian, uterine, cervical, prostate, liver, colon, pancreatic and colon vorsian, uterine, cervical, prostate, liver, colon, pancreatic and colon pain cancer. AAMO6005 to AAM1356 represent CYPIB1 peptides which can bind to human leukocyte antigens (HIAs), as well as other amino acid sequence used in the exemplification of the present invention
                            Disclosure; Page 69; 121pp; English
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Sequence 10 AA;

ö Gaps . Score 37; DB 4; Length 10; Pred. No. 23; 1; Mismatches 2; Indels 45.1%; 70.0%; Query Match
Best Local Similarity 70.00
7; Conservative

AAM12950 standard; peptide; 10 AA. RESULT 7

AAM12950; THE STANDARD STANDARD

(first entry) 09-OCT-2001

HLA-A26 decamer #87,

Cytochrome P450 1B1; CYP1B1; cancer; immunotherapy; diagnosis; HLA; human leukcoyte antigen; cytocxic T-lymphocyte; CTL; MHC; APC; major histocompatibility complex; antigen presenting cell; melanoma; lymphoma; carcinoma; multiple myeloma; leukaemia.

Homo sapiens.

WO200135810-A2

25-MAY-2001

15-NOV-2000; 2000WO-US031513.

99US-0165590P. LS-NOV-1999; (DAND) DANA FARBER CANCER INST INC. (UYBO-) UNIV BOSTON.

Vonderheide RH, Sherr D, Nadler LM, Maecker B; Von Bergwelt- Baildon M; Schultze JL,

WPI; 2001-355537/37.

Treating a cancer patient involves administering to patient a cytotoxic T-lymphocyte, an antigen presenting cell that activates T cells, a peptide of cytochrome P450 1B1, or a polynucleotide encoding the peptide.

```
The present invention describes a method for treating a patient having, or is at risk of having, a cell expressing cytochrome P450 1B1 (CYP1B1), which involves administering to the patient a cytocoxic T-lymphocyte (CTL) that kills the cell, an antigen presenting cell (APC) that cativates CTL, a peptide of CYP1B1 that binds to major histocompatibility complex (MMC) molecule, or a nucleic acid molecule encoding CYP1B1 or peptide of CYP1B1. Also described are: (1) a method for assessing the complex (MMC) molecule, involving measuring the level of CYP1B1 or peptide of CYP1B1 or the peptide of CYP1B1 or a peptide of CYP1B1 that binds to MHC complex molecule, involving measuring the level of CTL capecific for CYP1B1 or the peptide of CYP1B1 in a sample from the patient complex peptide of CYP1B1 in a sample from the patient complex peptide of MHC complex restricted fashion; and (4) an ex vivo generated APC (III) that specifically kills a cell expressing CYP1B1 in a wife complex molecule. The method is useful for treating a patient capecific of a MHC complex restricted fashion; and (4) and extend the method is useful for the prevention, treatment and diagnosis of cancer, e.g. melanoma, lymphoma, carcinoma, sarcoma, multiple myeloma, leukemia, and cludy ovarian, uterine, cervical, prostate, liver, colon, pancreatic and chain cancer. AAM06905 to AAM13566 represent CYP1B1 peptides which can bind to human leukcoyte antigens (HAMS), as well as other amino acid sequence used in the exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1; Mismatches
                      Disclosure; Page 89; 121pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity 70.00.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           6 FVSPRPLPVV 15
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AAM08833 standard; peptide; 10 AA. 09-OCT-2001 (first entry) HLA-Al decamer #12. AAM08833; AAM08833 A CONTRACTOR OF THE CONTRACT O

Cytochrome P450 1B1; CYP1B1; cancer; immunotherapy; diagnosis; HLA; human leukcoyte antigen; cytocxic T-lymphocyte; CT1, MHC; APC; major histocompatibility complex; antigen presenting cell; melanoma; lymphoma; carcinoma; multiple myeloma; leukaemia.

WO200135810-A2.

15-NOV-2000; 2000WO-US031513.

25-MAY-2001

(DAND) DANA FARBER CANCER INST INC. (UYBO-) UNIV BOSTON. 99US-0165590P. 15-NOV-1999;

Vonderheide RH, Sherr D, Nadler LM, Maecker B; Schultze JL, Vonderheid Von Bergwelt- Baildon M;

WPI; 2001-355537/37.

Treating a cancer patient involves administering to patient a cytotoxic T -lymphocyte, an antigen presenting cell that activates T cells, a peptide of cytochrome P450 1B1, or a polynucleotide encoding the peptide.

Disclosure; Page 68; 121pp; English

The present invention describes a method for treating a patient having, or is at risk of having, a cell expressing cytochrome P450 1B1 (CYPIB1), which will have be patient a cytocoxic T-lymphocyte (CTL) that kills the cell, an antigen presenting cell (APC) that activates CTL, a peptide of CYPIB1 that binds to major histocompatibility complex (MHC) molecule, or a nucleic acid molecule encoding CYPIB1 or peptide of CYPIB1 or method for assessing the complex (MHC) molecule, involving measuring the level of fimmunity of a patient to CYPIB1 or a peptide of CYPIB1 that binds to MHC complex molecule, involving measuring the level of CTL specific for CYPIB1 or the peptide of CYPIB1 in a sample from the patient compensated CTL (II) that binds to MHC complex molecule; (3) an exvivo generated APC (III) that specifically kills a cell expressing CYPIB1 in a specific of a MHC complex restricted feshion; and (4) an ex vivo generated APC (III) that presents a peptide of a CYPIB1 in the context of a MHC complex molecule. The method is useful for treating a patient complex molecule. The method is useful for treating a patient complex molecule. The method is useful for treating a patient complex molecule, or avoing a cell that expresses CYPIB1. The method is useful for the prevention, treatment and diagnosis of cancer, e.g. melanoma, lymphoma, carcinoma, sarcoma, multiple myeloma, leukemia, and compance in the exemplification of the present invention

Sequence 10 AA;

Gaps . 0 Score 37; DB 4; Length 10; Pred. No. 23; 2; Indels 1; Mismatches 45.1%; 70.0%; 7; Conservative Query Match Best Local Similarity Matches

; 0

6 FVSPRPLPVV 15 FLDPRPLTVV 10

RESULT

AAM09489 standard; peptide; 10 AA. AAM09489

09-OCT-2001 (first entry) AAM09489;

HLA-B *0702 decamer #115.

Cytochrome P450 1B1; CYP1B1; cancer; immunotherapy; diagnosis; HLA; human leukocyte antigen; cytocoxic T-lymphocyte; CTL, MHC; APC; madjor histocompatibility complex; antigen presenting cell; melanoma; lymphoma; carcinoma; sarcoma; multiple myeloma; leukaemia.

WO200135810-A2.

25-MAY-2001.

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15-NOV-2000; 2000WO-US031513.

99US-0165590P. 15-NOV-1999; (DAND) DANA FARBER CANCER INST INC. (UYBO-) UNIV BOSTON.

Nadler LM, Maecker B; Sherr D, Schultze JL, Vonderheide RH, Von Bergwelt- Baildon M;

WPI; 2001-355537/37.

Treating a cancer patient involves administering to patient a cytotoxic T-lymphocyte, an antigen presenting cell that activates T cells, a peptide of cytochrome P450 1B1, or a polynucleotide encoding the peptide.

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or is at risk of having, a cell expressing cytochrome P450 IBI (CYPIB1), which involves administering to the patient a cytochrome P450 IBI (CYPIB1), which involves administering to the patient a cytochrome P450 IBI (CYPIB1), that kills the cell, an antigen presenting cell (APC) that activates CTL, a peptide of CYPIBI that binds to major histocompatibility complex (MHC) molecule, or a nucleic acid molecule encoding CYPIBI or peptide of CYPIBI. Also described are: (1) a method for assessing the level of immunity of a patient to CYPIBI or a peptide of CYPIBI that binds to MHC complex molecule, involving measuring the level of CTL binds to MHC complex molecule, involving measuring the level of CTL crowled in the peptide of CYPIBI in a sample from the patient or (2) a CYPIBI peptide (I) that binds to MHC complex molecule; (3) an exvivo generated CTL (II) that specifically kills a cell expressing CYPIBI or a peptide of a CYPIBI in the context of a MHC complex molecule. The method is useful for treating a patient or service of having a cell that expresses CYPIBI. The method is useful for the prevention, treatment and diagnosis of cancer e.g. melanoma, lamphoma, carcinoma, sactoma, multiple myeloma, leukemia, and brain cancer. AAM06905 to AAM1356s represent CYPIBI peptides which can bind to human leukocyte antigens (HLMAS), as well as other amino acid context of sequence used in the exemplification of the present invention
                                                                        The present invention describes a method for treating a patient having
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 37; DB 4; Length 10;
Pred. No. 23;
1; Mismatches 2; Indels
                      Disclosure; Page 71; 121pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       45.1%;
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 10 AA;
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d
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AAW11111 standard; peptide; 13 AA. 25-JUN-1997 (first entry) 10 1 FLDPRPLTVV AAW11111; RESULT 10 AAW11111

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Src SH3 domain-binding peptide used in signal transduction modulation. Src; SH3; Src homology region 3; binding affinity; oncogenic protein; protein tyrosine kinase; signal transduction; RNA processing; 94US-00278865. 95US-00483555. 95WO-US009382 trafficking, translation. WO9603649-A1. 24-JUL-1995; 22-JUL-1994; 07-JUN-1995; 08-FEB-1996. Synthetic.

Der CJ; Quilliam LA, Thorn JM, (UYNC-) UNIV NORTH CAROLINA. WPI; 1996-117151/12. Sparks AB, Kay BK,

Peptide with binding affinity for Src homology region 3 (SH3) domains of proteins - useful for e.g. modulating signal transduction pathways at the cellular level, esp. protein tyrosine kinase-mediated.

Claim 38; Page 87; 116pp; English

· 0

Gaps

0

42.7%; Score 35; DB 2; Length 13; 70.0%; Pred. No. 65; 3; Indels ive 0; Mismatches 3; Indels

Ouery Match
Best Local Similarity 70...
7, Conservative

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AAWI1098-WI1124 are peptides that bind to the Src SH3 domain. The SH3 binding peptides are useful in modulating signal transduction pathways at the cellular level (especially protein tyrosine kinase-mediated), modulating oncogenic protein activity, or providing compounds for the development of drugs with the ability to modulate broad classes, as well as specific classes, of proteins involved in signal transduction and also
                                                                            for regulating the processing, trafficking or translation of RNA. Conjugates of the peptides with detectable labels or imaging agents are useful for imaging cells, tissues and organs in which Src or Src-related
                                                                                                                                                                                                                                                                                                                                                                                               domain; SH3 domain; oncogene; consensus; lambda 14;
                                                                                                                                                             DB 2; Length 13;
65;
                                                                                                                                                                                  0; Indels
                                                                                                                                                           Score 35; DB 2
Pred. No. 65;
3; Mismatches
                                                                                                                                                                                                                                                                                                                                                                       Proline-rich SH3 binding peptide L14-2.
                                                                                                                                                                                                                                                                                       AAW05098 standard; peptide; 13 AA.
                                                                                                                                                             42.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     94US-00230047.
                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                    5; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                           random display library.
                                                                                                                 proteins are expressed
                                                                                                                                                                                                                                                                                                                                        (revised)
                                                                                                                                                                                                          8 SPRPLPVV 15
                                                                                                                                                                                                                                 2 TPRPLPLI 9
                                                                                                                                                           Query Match
Best Local Similarity
Matches 5; Conserv
                                                                                                                                       Sequence 13 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    19-APR-1994;
                                                                                                                                                                                                                                                                                                                                                                                               Src homology
                                                                                                                                                                                                                                                                                                                                      25-MAR-2003
17-DEC-1996
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                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
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                                                                                                                                                                                                                                                                  RESULT 11
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X000000000000X8
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0; Gaps

A fusion protein of glutathione-S-transferase, protein kinase A and Src SH3 domain was used to screen a mouse embryonal cDNA library in lambda EXIOX. Three positive clones were identified and were designated Li1, Li4 and Li35. The insert from clone Li4 contains a 221 bp region with approximately 90% identity to an anonymous cDNA (EST06380) directionally cloned from a human infant brain cDNA library. The deduced amino acid sequence contains two proline-rich, putative SH3-binding sequences. The present peptide sequence is derived from the putative SH3-binding site at residues 241-253 of Li4 and was used in an assay to determine binding to the SH3 domains of c-src and other proteins. (Updated on 25-MAR-2003 to New DNA encoding proteins that bind to Src-homology domain 3 - used to inhibit signal transduction, e.g. in cases of T cell activation. Example 1; Col 16; 39pp; English. correct PI field.)

Cheadle C, South VJ,

Ricca GA,

Ivashchenko YD,

French SM,

Searfoss GH;

WPI; 1996-361965/36.

(RHON) RHONE POULENC RORER PHARM INC.

94US-00230047.

Sequence 13 AA;

9 PRPLPV 14

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cells, e.g. responsible for neoplastic conditions. Specific examples of Src proteins with SH3 domains are those known by the abbreviations p85, Fyn, Lyn, Hck, Syk, GrbJ, Gap, STAT, p47-phox, p67-phox, and Brk. The present sequence represents a nonamer peptide used in an example from the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    spiro[Pyrrolidine-2':3-piperidin-2-ones] and their fused ring analogues used, e.g. to modulate signal transduction protein interactions, inhibit cell growth and for use in treating infections and cancer.
                                                                                                                                                                   Src howology 3 domain; SH3 domain; peptidomimetic; mimetic; modulator; beta-turn template; spirolPyrrolidine-2':3-piperidin-2-onee]; fused ring analogue; signal transduction protein interaction; inhibit cell growth; infection; cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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                                                                                                                                   Src homology 3 domain binding nonamer peptide #1.
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100.0%; Pred. No. 1.4e+06;
ative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                                                                                                                                                             /note= "acetylated"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 12; Page 59; 73pp; English.
                                  AAW94619 standard; peptide; 9 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                  98WO-US010861.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                97US-00864241.
98US-0085889P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Castelhano AL, Witter DJ;
                                                                                                29-APR-1999 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
6; Conserve
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1999-080804/07.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 9 AA;
                                                                                                                                                                                                                                                                                                             Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                  28-MAY-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                03-DEC-1998.
                                                                                                                                                                                                                                                           Synthetic.
                                                                  AAW94619;
RESULT 12
                 AAW94619
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1 AFLDPRPLTV 10

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Cytochrome P450 1B1; CYP1B1; cancer; immunotherapy; diagnosis; HLA; human leukcoyte antigen; cytochic T-lymphocyte; CTL; MHC; APC; humaior histocompatibility complex; antigen presenting cell; melanoma; lymphoma; carcinoma; sarcoma; multiple myeloma; leukaemia.
                                                                                                                                                                                        FARBER CANCER INST INC. BOSTON.
              AAM09403 standard; peptide; 10 AA.
                                                                                                                                                                                                              Schultze JL, Vonderheide RH,
Von Bergwelt- Baildon M;
                                                                                                                                                           L5-NOV-2000; 2000WO-US031513.
                                                                                                                                                                          99US-0165590P.
                                            09-OCT-2001 (first entry)
                                                                                                                                                                                                                                     WPI; 2001-355537/37.
                                                           HLA-Al decamer #122.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 10 AA;
                                                                                                                              WO200135810-A2.
                                                                                                               Homo sapiens.
                                                                                                                                                                                       (DAND ) DANA
(UYBO-) UNIV
                                                                                                                                                                          15-NOV-1999;
                                                                                                                                             25-MAY-2001.
                            AAM09403;
RESULT 13
       AAM09403
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Maecker B;

Sherr D, Nadler LM,

Query Match Best Local Similarity

Conservative

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41.5%; Score 34; DB 4; Length 10; 60.0%; Pred. No. 72; 2; Indels tive 2; Mismatches 2; Indels
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The present invention describes a method for treating a patient having, or is at risk of having, a cell expressing cytochrome P450 IB1 (CYPIB1), which involves administering to the patient a cytocoxic T-lymphocyte (CTL) that kills the cell, an antigen presenting cell (APC) that activates CTL, a peptide of CYPIB1 that binds to major histocompatibility activates CTL, nolecule are (1) a method for assessing the complex (MMC) molecule are: (1) a method for assessing the competide of CYPIB1. Also described are: (1) a method for assessing the level of immunity of a patient to CYPIB1 or a peptide of CYPIB1 that binds to MMC complex molecule, involving measuring the level of CTL specific for CYPIB1 or the peptide of CYPIB1 in a sample from the patient (2) a CYPIB1 peptide (I) that binds to MMC complex molecule, (3) an exvivo generated APC (III) that specifically kills a cell expressing CYPIB1 in a specific of a MHC complex-restricted fashion; and (4) and extrement of a method is useful for treating a patient complex molecule. The method is useful for treating a patient complex molecule. The method is useful for treating a patient complex molecule. The method is useful for treating a patient of a wing or is at risk of having a cell that expresses CYPIB1. The method is useful for the prevention, treatment and diagnosis of cancer. Graving convarian, uterine, cervical, prostate, liver, colon, pancreatic and colon, pancreatic and colon vancreatic and to brain cancer. AAMO6905 to AAM13506 represent CYPIB1 peptides which can be brain cancer used in the exemplification of the present invention
Treating a cancer patient involves administering to patient a cytotoxic T-lymphocyte, an antigen presenting cell that activates T cells, a peptide of cytochrome P450 1B1, or a polynucleotide encoding the peptide.
                                                                                                                                                                                                                                                                                    Disclosure; Page 71; 121pp; English.
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The present invention describes a method for treating a patient having, or is at risk of having, a cell expressing cytochrome P450 1B1 (CYPIB1), which involves administering to the patient a cytocoxic T-lymphocyte (CTL) that kills the cell, an antiquen presenting cell (APC) that activates CTL, a peptide of CYPIB1 that binds to major histocompatibility complex (MHC) molecule, or a nucleic acid molecule encoding CYPIB1 or peptide of CYPIB1. Also described are: (1) a method for assessing the complex molecule, involving measuring the level of CYPIB1 to the peptide of CYPIB1 in a sample from the patient or specific for CYPIB1 or the peptide of CYPIB1 in a sample from the patient or vivo generated CTL (II) that binds to MHC complex molecule, (3) an exvivo generated CTL (II) that specifically kills a cell expressing CYPIB1 in a specific of a MHC complex-restricted fashion; and (4) an ex vivo generated APC (III) that presents a peptide of a CYPIB1 in the context of a MHC complex molecule. The method is useful for treating a patient construct the prevention, treatment and diagnosis of cancer. Grancer and the context of melanoma, lymphoma, carcinoma, sarcoma, multiple myeloma, leukemia, and cill gone are an expensive and context of melanoma, lymphoma, carcinoma, sarcoma, multiple myeloma, leukemia, and context of the prevention, treatment and diagnosis of cancer. ApM06905 to AAM1356 represent CYPIB1 peptides which can brain cancer. Aam06905 to AAM1356 represent CYPIB1 peptides which can brain cancer and the exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       a peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Treating a cancer patient involves administering to patient a cytotoxic -lymphocyte, an antigen presenting cell that activates T cells, a peptid of cytochrome P450 1B1, or a polynucleotide encoding the peptide.
                                                                                                                                                                                                                Cytochrome P450 1B1; CYP1B1; cancer; immunotherapy; diagnosis; HLA; human leukcoyte antigen; cytocoxic T-lymphocyte; CTL; MHC; APC; major histocompatibility complex; antigen presenting cell; melanoma; lymphoma; carcinoma; sarcoma; multiple myeloma; leukeemia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Maecker B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         41.5%; Score 34; DB 4; Length 10; 60.0%; Pred. No. 72;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Nadler LM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pred. No. 72;
2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Vonderheide RH, Sherr D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; Page 93; 121pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (DAND ) DANA FARBER CANCER INST INC.
                                           AAM13226 standard; peptide; 10 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          99US-0165590P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                               15-NOV-2000; 2000WO-US031513.
                                                                                                                           09-OCT-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity 60.0
nes 6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Von Bergwelt- Baildon M;
                                                                                                                                                                            HLA-A26 decamer #289.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2001-355537/37.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (UYBO-) UNIV BOSTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 10 AA;
                                                                                                                                                                                                                                                                                                                                                                          WO200135810-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Schultze JL,
                                                                                                                                                                                                                                                                                                                                 Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        15-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                        25-MAY-2001.
                                                                                    AAM13226;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
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Matches
RESULT 14
                      AAM13226
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Gaps

. 0

5 SFVSPRPLPV 14

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Gaps

·.

2; Indels

5 SFVSPRPLPV 14

1 AFLDPRPLTV 10

RESULT 15 AAM11873

AAM11873 standard; peptide; 10 AA.

AAM11873;

09-OCT-2001 (first entry)

HLA-A *0202 decamer #110.

Cytochrome P450 1B1, CYP1B1; cancer; immunotherapy; diagnosis; HLA; human leukcoyte antigen, cytotoxic T-lymphocyte; CTL; MHC; APC; major histocompatibility complex; antigen presenting cell; melanoma; lymphoma; carcinoma; sarcoma; multiple myeloma; leukaemia.

Homo sapiens.

WÓ200135810-A2.

25-MAY-2001,

15-NOV-2000; 2000WO-US031513.

99US-0165590P. 15-NOV-1999;

(DAND) DANA FARBER CANCER INST INC. (UYBO-) UNIV BOSTON.

Maecker B; Nadler LM, Sherr D, Schultze JL, Vonderheide RH, Von Bergwelt- Baildon M;

WPI; 2001-355537/37.

Treating a cancer patient involves administering to patient a cytotoxic T-lymphocyte, an antigen presenting cell that activates T cells, a peptide of cytochrome P450 1B1, or a polynucleotide encoding the peptide.

Disclosure; Page 84; 121pp; English.

The present invention describes a method for treating a patient having, or is at risk of having, a cell expressing cytochrome P450 1B1 (CYPIB1), which involves administering to the patient a cytocoxic T-lymphocyte (CTL) that kills the cell, an antigen presenting cell (APC) that activates CTL, a peptide of CYPIB1 that binds to major histocompatibility complex (MMC) molecule, or a nucleic acid molecule encoding CYPIB1 or peptide of CYPIB1 Also described are: (1) a method for assessing the cell immunity of a patient to CYPIB1 or a peptide of CYPIB1 that binds to MMC complex molecule, involving measuring the level of CYPIB1 peptide (1) that binds to MMC complex molecule, involving measuring the level of CYPIB1 corporated (TPIB1) a sample from the patient (1) (2) a CYPIB1 peptide (1) that binds to MMC complex molecule, (3) an exvivo generated APC (III) that specifically kills a cell expressing CYPIB1 in a specific of a MMC complex molecule. The method is useful for treating a patient and Complex molecule. The method is useful for treating a patient context of a MMC complex molecule. The method is useful for treating a patient context of a wing or is a trisk of having a cell that expresses CYPIB1 in the context of melanoma, lymphoma, carcinoma, sarcoma, multiple myeloma, leukemia, and cis useful for the prevention, treatment and diagnosis of cancer, e.g. melanoma, lymphoma, carcinoma, sarcoma, multiple myeloma, leukemia, and chain cancer. AAMMG505 to AAMM1356 represent CYPIB1 peptides which can bind to human leukocyte antigens (HLAs), as well as other amino acid sequence used in the exemplification of the present invention

Sequence 10 AA;

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Gaps
                                                  ;
0
Query Match
Best Local Similarity 60.0%; Pred. No. 72;
Matches 6; Conservative 2; Mismatches 2; Indels
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. 0

1 AFLDPRPLTV 10

Search completed: April 2, 2004, 10:08:42 Job time : 46 secs

5 SFVSPRPLPV 14

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GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
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OM protein - protein search, using sw model

2, 2004, 09:58:38; Search time 7.66667 Seconds (without alignments) 101.876 Million cell updates/sec April Run on:

US-10-009-049-1 84

1 CRAHSFGSPRPLPVV 15 Perfect score: Sequence:

BLOSUM62 Scoring table:

141681 seqs, 52070155 residues Gapop 10.0 , Gapext 0.5 Searched:

801 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 15

Maximum Match 100% Post-processing: Minimum Match 0%

Listing first 45 summaries

SwissProt_42:* Database :

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

CITAMADITE

15 AA.

STANDARD;

(Rel. 34, Created) (Rel. 34, Last sequence update)

01-OCT-1996 01-OCT-1996 UC29 MAIZE

P80635; UC29 MAIZE ID UC29 MA AC P80635, DT 01-OCT-DT 01-OCT-

P28499 oncorhynchu P41333 scyliorhinu	P80621 zea mays (m P81817 carcinus ma	P81818 carcinus ma P81820 carcinus ma	P81821 carcinus ma P81780 herpes simp	P81822 carcinus ma P18523 manduda sex		P56919 rana tempor
TKNA_ONCMY TKNA_SCYCA	UC15 MAIZE AL14 CARMA	AL15_CARMA AL17_CARMA	AL18_CARMA VGLG_HSV2B	AL19_CARMA FARP_MANSE	PNEU RAT	TEMD_RANTE
н н	пп.	7 7		н -	п	7
11	14 5	00 00	∞ ∞	10	10	13
22.6	22.6	21.4	21.4	21.4	21.4	21.4
919	61	18 18	18 18	18	18	18
3.4 3.5	36	80 6	4 4 1 1	4. 4. 2. E.	44	45

ALIGNMENTS

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0
DEPTI BOTJA STANDARD; PRT; 13 AA.

(C P01020; P30421;
(C P01020; P30421;
(C L1986 (Rel. 01, Created)
(C LTEB-1994 (Rel. 28, Last sequence update)
(C LTEB-1994 (Rel. 41, Last annotation update)
(C SPEP-2003 (Rel. 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SPECIES=B.insularis, TISSUB=Venom,
MEDLINE=90351557; PubMed=2386615;
Cintra A.C.O., Vieira C.A., Gigalio J.R.;
Cintra A.C.O., Vieira C.A., Gigalio J.R.;
Special activity of bradykinin potentiating peptides from Bothrops insularis snake venom.";
J. FUNCTION: This peptide both inhibits the activity of the angiotensin-converting enzyme and enhances the action of bradykinin by inhibiting the kinases that inactivate it.

II acts as an indirect hypotensive agent.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    "Anglotensin-converting enzyme inhibitors from the venom of Bothrops jararaca. Isolation, elucidation of structure, and synthesis."; Biochemistry 10:4033-4039(1971).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDLINE=72118526; PubMed=4334402;
Ondetti M.A., Williams N.J., Sabo E.F., Pluscec J., Weaver E.R.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            .,
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Hypotensive agent; Pyrrolidone carboxylic acid.

MOD RES 1 1 PYRROLIDONE CARBOXYLIC ACID.

SEQÜENCE 13 AA; 1388 MW; 6824FC97D83D6774 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                31.0%; Score 26; DB 1; Length 13; 55.6%; Pred. No. 2.38+02; tive 1; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SPECIES=B.jararaca; TISSUE=Venom;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        5; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               7 GSPRPLPVV 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GWPRPGPEI 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity
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                                             임
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                                                                                                                                                                        Touzet P., Riccardi F., Morin C., Damerval C., Huet J.-C.,
Pernollet J.-C., Zivy M., de Vienne D.;
"The maize two dimensional gel protein database: towards an integrated
                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Electrophoresis 20:1098-1108(1999).
-1- FUNCTION: May be involved in the regulation of photosystem II.
-1- SUBCELLULAR LOCATION: Chloroplast thylakoid membrane; associated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        15-JUL-1999 (Rel. 38, Created)
15-JUL-1999 (Rel. 38, Last sequence update)
15-MAR-2004 (Rel. 43, Last annotation update)
0xygen-evolving enhancer protein 2 (ORE2) (23 kDa subunit of oxygen evolving system of photosystem II) (Fragment)
                                                                                                                                                                                                                   genome analysis program.";
Theor. Appl. Genet. 93:997-1005(1996).
-!- MISCELLANBOUS: On the 2D-gel the determined pI of this unknown protein is: 4.9, its MW is: 37.6 kDa.
Maize-2DPAGE; P80635; COLEOPTILE.
                                                          Eukaryota, Viridiplantae, Streptophyta, Embryophyta, Tracheophyta,
Spermatophyta, Magnoliophyta, Liliopsida, Poales, Poaceae,
PACCAD clade, Panicoideae, Andropogoneae, Zea.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota, Viridiplantae, Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Coniferopsida; Coniferales; Pinaceae; Pinus.
NCBI_TaxID=71647;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             with the photosystem II complex (By similarity).
-!- MISCELLANEOUS: On the 2D-gel the determined pI of this protein (spot N179) is: 5.9, its MW is: 22 kDa.
-!- SIMILARITY: Belongs to the psbP family.
Photosynthesis; Photosystem II; Chloroplast; Thylakoid; Membrane.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDLINE=99274088; PubMed=10344291;
Costa P., Pionneau C., Bauw G., Dubos C., Bahrman N., Kremer A.,
Frigerio J.-M., Plomion C.;
"Separation and characterization of needle and xylem maritime pine
                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
15-WAR-2004 (Rel. 43, Last annotation update)
Unknown protein from 2D-page of etiolated coleoptile (Spot 45)
                                                                                                                                                                                                                                                                                                                                                                                Score 26; DB 1; Length 15; Pred. No. 2.7e+02; 3; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            29.8%; Score 25; DB 1; Length 13; 50.0%; Pred. No. 3.3e+02; tive 3; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                   15 15 AA, 1679 MW; 3D53086B16018BC1 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            13 AA; 1294 MW; C6772B0D54D7C44D CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               13 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                 3; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pinus pinaster (Maritíme pine).
                                                                                                                                                                                                                                                                                                                                                                                  31.0%;
                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STANDARD;
                                                                                                                                                                                                                                                                                                     MaizeDB; 123960; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                               8 SPRPLPV 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  :|:|:
1 NPNPVPI 7
                                                                                                                                                           TISSUE=Coleoptile;
                                              Zea mays (Maize).
                                                                                                          NCBI_TaxID=4577;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TISSUE=Needle;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PSBP PINPS
P81668;
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                                   Fragment)
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NON TER
SEQUENCE
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g
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3 AHSFGSPR 10 |: ||:| 6 ANVFGAPK 13

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Cintra A.C.O., Vieland C.A., Giglio J.R.,

"Primary structure and biological activity of bradykinin potentiating peptides from Bothrops insularis snake venom.";

J. Protein Chem. 9:221-227(1990).

-!- FUNCTION: This peptide both inhibits the activity of the angiotensin-converting enzyme and enhances the action of bradykinin by inhibiting the kinases that inactivate it.

It acts as an indirect hypotensive agent.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cintra A.C.O., Vielra C.A., Giglio J.R.,
"Primary structure and biological activity of bradykinin potentiating peptides from Bothrops insularis snake venom.",
J. Protein Chem. 9:22-22/11990.
-!- FUNCTION: This peptide both inhibits the activity of the angiotensin-converting enzyme and enhances the action of bradykinin by inhibiting the kinases that inactivate it.
It acts as an indirect hypotensive agent.
PIR: D37196, D37196.
Hypotensive agent; Pyrrolidone carboxylic acid.
MOD RES
1 1 PYRROLIDONE CARBOXYLIC ACID.
SEQÜENCE 11 AA; 1143 MW; 20BBBF13C7741777 CRC64;
                                                                                                                  Bradykinin-potentiating peptide S4,3,2 (10C) (Angiotensin-converting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                   Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Lepidosauria, Squamata, Scleroglossa, Serpentes, Colubroidea,
Viperidae, Crotalinae, Bothrops.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Lepidosauria; Squamata; Scleroglossa; Serpentes; Colubroidea;
Viperidae; Crotalinae; Bothrops.
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0
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01-FRB-1994 (Rel. 28, Last sequence update)
28-FBB-2003 (Rel. 48, Last amontation update)
Bradykinin-potentiating peptide 54,1,2 (Angiotensin-converting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hypotensive agent; Pyrrolidone carboxylic acid.
MOD_RES 1 1 PYRROLIDONE CARBOXYLIC ACID.
SEQÜENCE 11 AA; 1199 MW; 20B25813C7741777 CRC64;
                                                                                                                                                   Bothrops insularis (Island jararaca) (Queimada jararaca).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Bothrops insularis (Island jararaca) (Queimada jararaca)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 24; DB 1;
Pred. No. 4e+02;
                                                                                (Rel. 28, Last sequence update) (Rel. 41, Last annotation update)
                                  11 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                  PRT;
                                                                                                                                                                                                                                                                                         MEDLINE=90351557; PubMed=2386615;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDLINE=90351557; PubMed=2386615;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ,
                                                                  (Rel. 25, Created)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  28.6%;
80.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                  STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  PIR; C37196; C37196.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
nes 4; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      7 GSPRP 11
                                                                                                                                  enzyme inhibitor)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       3 GPPRP 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                enzyme inhibitor)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NCBI_TaxID=8723;
                                                                                                                                                                                                                                                                             TISSUE=Venom;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TISSUE=Venom;
                                                                01-APR-1993
                                                                                  01-FEB-1994
28-FEB-2003
                                BPP3 BOTIN
P30423;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BPP4 BOTIN
P30424;
                                                                                                                                                                                                                                                         SECUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
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RESULT 4
BPP3 BOTIN
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-!- FUNCTION: Antibacterial peptide active against Gram-negative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Thernyeh S., Cociancich S., Briand J.-P., Hetru C., Bulet P., "The inducible antibacterial peptides of the hemipteran insect Palomena prasaina: identification of a unique family of proline-rich peptides and of a novel insect defensin.";
J. Insect Physiol. 42:81-89(1996).
-!- FUNCTION: Antibacterial peptide active against Gram-negative
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                                                                                                                                                                                                                                                                        Palomena prasina (Green shield bug).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Paraneoptera; Hemiptera; Euhemiptera; Heteroptera;
Panheteroptera; Pentatomomorpha; Pentatomoidea; Pentatomidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota, Metazoa, Arthropoda, Hěxapoda, Insecta, Pterygota,
Neoptera, Paraneoptera, Hemiptera, Euhemiptera, Heteroptera,
Panheteroptera, Pentatomomorpha, Pentatomoidea, Pentatomidae,
   Score 24; DB 1; Length 11;
Pred. No. 4e+02;
0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    28.6%; Score 24; DB 1; Length 15; 80.0%; Pred. No. 5.5e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Antibiotic; Insect immunity.
SEQUENCE 15 AA; 1838 MW; 21407E663CE46299 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     bacteria.
-!- INDUCTION: By bacterial infection.
Antibiotic; Insect immunity.
SEQUENCE 15 AA; 1893 MW; 23835D063B946299 CRC64;
                                                                                                                                                                                                            01-NOV-1995 (Rel. 32, Created)
01-NOV-1995 (Rel. 32, Last sequence update)
28-FBB-2003 (Rel. 41, Last annotation update)
Metalnikowin I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-NOV-1995 (Rel. 32, Created)
01-NOV-1995 (Rel. 32, Last sequence update)
28-FEB-2003 (Rel. 41, Last annotation update)
                                                                                                                                                                                 15 AA.
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Palomena prasina (Green shield bug).
     28.6%;
80.0%;
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                                   4; Conservative
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Best Local Similarity
Query Match
Best Local Similarity
                                                              7 GSPRP 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TISSUE=Hemolymph;
                                                                                                                                                                                                                                                                                                                                                           NCBI_TaxID=55431;
                                                                                                                                                                                                                                                                                                                                                                                                           IISSUE=Hemolymph;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 8 PRPRP 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NCBI_TaxID=55431;
                                                                                               3 GPPRP 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      bacteria
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28-FEB-2003
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P80409;
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P80408;
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                                   Matches
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                                                                                                                                                                                                                                                           Ascaris suum (Pig roundworm) (Ascaris lumbricoides).
Eukaryota, Metazoa, Nematoda, Chromadorea, Ascaridida, Ascaridoidea,
Ascarididae, Ascaris.
                                                                                                                                                                                                                                                                                                                                                                       Cowden C., Stretton A.O.W.;
"Eight novel FMRFamide-like neuropeptides isolated from the nematode
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 -! - FUNCTION: May be involved in the regulation of photosystem II.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           01-0CT-1996 (Rel. 34, Created)
01-0CT-1996 (Rel. 34, Last sequence update)
16-0CT-2001 (Rel. 40, Last annotation update)
06-0CT-2001 (Rel. 40, Last annotation update)
06xgen-evolving enhancer protein 2 (OBE2) (24 kDa subunit of oxygen evolving system of photosystem II) (Fragment).
Physocmitrella patens (Moss).
Bukaryota; Viridiplantes, Streptophyta; Embryophyta; Bryophyta;
Bryopsida; Funariidae; Funariales; Funariaceae; Physcomitrella.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TISSUE=Protonema;
MEDLINE=97275459; PubMed=9129336;
Kasten B., Buck F., Nuske J., Reski R.;
"Cytokinin affects nuclear- and plastome-encoded energy-converting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       -!- SIMILARITY: Belongs to the psbP family.
Photosynthesis; Photosystem II; Chloroplast; Thylakoid; Membrane;
Multigene family.
NON_TER 12 12
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-!- SIMILARITY: Belongs to the FARP (FMRFamide related peptide)
             Score 24; DB 1; Length 15;
Pred. No. 5.5e+02;
0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   27.4%; Score 23; DB 1; Length 9; 100.0%; Pred. No. 1.4e+05; ative 0; Mismatches 0; Indels
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SEQÜENCE 9 AA; 1012 MW; 524F073774176877 CRC64;
                                                                                                                                                                                                                    (Rel. 32, Last sequence update) (Rel. 33, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     12 AA.
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                                                                                                                                                                                                                                                                                                                                                       MEDLINE=95380362; PubMed=7651904;
                                                                                                                                                                                                                                                   FMRFamide-like neuropeptide AF9.
                                                                                                                                                                                                         01-NOV-1995 (Rel. 32, Created)
01-NOV-1995 (Rel. 32, Last seq
             28.6%;
80.0%;
Query Match
Best Local Similarity 80.v.
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Best Local Similarity luv...
4; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     plastid enzymes.";
Planta 201:261-272(1997).
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Neuropeptide; Amidation.
MOD RES 9 9
                                                                                                                                                                            STANDARD;
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                                                                        9 PRPLP 13
                                                                                                    8 PRPWP 12
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                                                                                                                                                                                                                                                                                                              NCBI_TaxID=6253;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         4 PRPL 7
                                                                                                                                                                                                                                                                                                                                                                                                    Ascaris suum."
                                                                                                                                                                                                                                       01-FEB-1996
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NCBI_TaxID=6233
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P82155:
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ALL4_CYDPO
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Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha; Oestroidea;
Calliphoridae; Calliphora.
                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
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Proc. Natl. Acad. Sci. U.S.A. 90:2456-2460(1993).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Duve H., Thorpe \lambda.; "Distribution and functional significance of Leu-callatostatins in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDLINE=93211980; PubMed=8460157;
Duve H., Johnsen A.H., Scott A.G., Yu C.G., Yagi K.J., Tobe S.S.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Duve H., Johnsen A.H., Maestro J.-L., Scott A.G., Winstanley Davey M., Bast P.D., Thorpe A.;
"Lepidopteran peptides the allatostatin superfamily.";
Peptides 18:1301-1309(1997),
-! SIMILARITY: Belongs to the allatostatin family.
                                       Score 23; DB 1; Length 12; Pred. No. 6.4e+02;
                                                                                 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  26.2%; Score 22; DB 1; Length 8; 66.7%; Pred. No. 1.4e+05;
  8D2B0D54D7C44DC5 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        8 AA; 926 MW; C82879D5AB477415 CRC64;
                                                                                                                                                                                                                                                                                                                                    30-MAY-2000 (Rel. 39, Last sequence update) 30-MAY-2000 (Rel. 39, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  01-NOV-1995 (Rel. 32, Created)
01-NOV-1995 (Rel. 32, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation update)
Callarostatin 4 (Leu-callatostatin 4).
                                                                                                                                                                                                                                                                         8 AA.
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                                                                                 2; Mismatches
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                                                                                                                                                                                                                                                                         PRT;
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MEDLINE=94291167; PubMed=8020069;
                                                                                                                                                                                                                                                                                                                                                                                               Cydia pomonella (Codling moth).
                                                                                                                                                                                                                                                                                                            30-MAY-2000 (Rel. 39, Created)
                                         27.4%;
57.1%;
12 AA; 1182 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TISSUE=Thoracic ganglion;
                                                                                 4; Conservative
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                                                                                                                                                                                                                                                                       STANDARD;
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6 ANVFGAP 12
                                                                                                                          3 AHSFGSP 9
                                                           Best Local Similarity
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RPYSFG 7
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                                                                                                                                                                                                                                                                                                                                                                            Cydiastatin 3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TISSUE=Larva;
                                                                                                                                                                                                                                                                  ALL3 CYDPO
P82154;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CALVO
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SEQUENCE
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                                         Query Match
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P41840;
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ALL4 CALVO
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                                                                                 Matches
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the blowfly Calliphora vomitoria.";
Call Tissue Res. 276:367-379(1994).
-!- FUNCTION: May act as a neurotransmitter or neuromodulator and play a role in the integration of information within the brain. May be involved in the control of visceral muscles due to its ability to behave as potent inhibitors of peristalic movements. May also fulfill a neurohormonal role on muscles of the gut and heart.
-!- TISSUE SPECIFICITY: Brain, suboseophageal ganglion, retrocerebral complex. Thoracico-abdominal ganglion, peripheral neurosecretory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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Duve H., Johnsen A.H., Maestro J.-L., Scott A.G., Winstanley D., Davey M., East P.D., Thorpe A.;
"Lepidopteran peptides of the allatostatin superfamily.";
Peptides 18:1301-1309(1997).
-!- SIMILARITY: Belongs to the allatostatin family.
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                                                                                                                                                                                                                                                                                                                                                                                                                              Length 8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1; Indels
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Panagrolalmoidea, Panagrolaimidae, Panagrellus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                16-OCT-2001 (Rel. 40, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
FMRFamide-like neuropeptide PF6 (NGAPQPFVRF-amide).
                                                                                                                                                                                                                   system and integrine.
-!- SIMILARITY: Belongs to the allatostatin family.
PIR; D47393; D47393.
                                                                                                                                                                                                                                                                                                                                                                          8 AA; 954 MW; D32879D5AB47740A CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    8 8 AMIDATION.
8 AA; 910 MW; 922879D5AB47740D CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                           26.2%; Score 22; DB 1; L 66.7%; Pred. No. 1.4e+05; tive 1; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  30-WAY-2000 (Rel. 39, Created)
30-WAY-2000 (Rel. 39, Last sequence update)
30-WAY-2000 (Rel. 39, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        B AA.
                                                                                                                                                                                                                                                                                                                              AMIDATION.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Cydia pomonella (Codling moth).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         16-OCT-2001 (Rel. 40, Created)
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity 66.7
Matches 4; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               4; Conservative
                                                                                                                                                                                                                                                                                                   Neuropeptide; Amidation.
MOD_RES 8 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STANDARD;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Panagrellus redivivus.
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2 RPYSFG 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2 RAHSFG 7
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SEQUENCE
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-!- FTM: S-linked glycan consists of GlC-GlC-GlC trisaccharide.
-!- MISCELLANEOUS: The identity of the glycoprotein from which this peptide is derived is unknown. No physiological function has been attributed.
          Moffett C.L., Marks N.J., Halton D.W., Thomson D.P., Geary T.G., Molectt C.L., Marks N.J., Halton D.W., Thomson D.P., Geary T.G., Maule A.G.,
"Isolation, characterization and pharmacology of RWRFamide-related peptides (FaRFs) from free-living nematode, Panagrellus redivivus.";
Submitted (JUL-2000) to Swiss-Prot.
-! FUNCTION: Myoactive.
-! - FUNCTION: Myoactive.
                                                                                                                                                                                                                                                              0; Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                    21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last sequence update)
16-OCT-2001 (Rel. 40, Last annocation update)
Erythrocyte membrane glycopeptide.
Homo sapiens (Human).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi;
Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tigerinin-4.

Hoplobatrachus tigerinus (Indian bull frog) (Rana tigerina).

Bukaryota, Metazoa, Chordata, Craniata, Verrebrata, Euteleostomi,

Amphibia, Batrachia, Anura, Neobatrachia, Ranoidea, Ranidae,

Hoplobatrachus
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                                                                                                                                                                                                                            Score 22; DB 1; Length 10; Pred. No. 7.6e+02;
                                                                                                                                                                                                                                                              0; Indels
                                                                                                                                                                             AMIDATION.
CB13E4C9D776C76D CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 1 S-LINKED (GLC. .).
10 AA; 1049 MW; 2339BFEEAAIFSBIEB CRC64;
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16-OCT-2001 (Rel. 40, Created)
16-OCT-2001 (Rel. 40, Last sequence update)
10-OCT-2003 (Rel. 42, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                           10 AA.
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                                                                                                                                                                                                                              26.2%;
60.0%;
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SEQUENCE 10 AA; 1132 MW;
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Best Local Similarity 60.v.
Sectorative 3; Conservative
                                                                                                                                                             Neuropeptide; Amidation.
MOD_RES 10 10
                                                                                                                                                                                                                                                                                                                                                                                                           STANDARD;
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les 3; Conserv
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GLEM_HUMAN
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Gaps
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Devi A.S., Nagaraj R., Sitaram N.;
"Tigerinins: novel antimicrobial peptides from the Indian frog Rana
                                                                                                                                                                                                                                                                                                                                   tigerina.";
J. Biol. Chem. 276:2701-2707(2001).
J. Biol. Chem. 276:2701-2707(2001).
J. Biol. Chem. 276:2701-2707(2001).
J. Shorton, M. Libacterial activity against B. subtilis, E. coli, S. aureus, M. Luteus, P. putida and S. cerevisiae.
J. SUBELLUTAR LOCATION: Secreted.
J. TISSUB SPECTROMETRY: NA.1.
J. MASS SPECTROMETRY: WH=1247; METHOD=MALDI.
Amphibian defense peptide; Antibiotic.
DISULFID
J. L. L. AMPHIBIAN ANTIBLE A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ..
                               SEQUENCE, FUNCTION, MASS SPECTROMETRY, AND DISULFIDE BONDS.
TISSUE=Skin secretion;
PubMed=11031261;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25.0%; Score 21; DB 1; Length 11; 75.0%; Pred. No. 1.2e+03; tive 1; Mismatches 0; Indele
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE 11 AA; 1248 MW; 117D8EFD37605DCB CRC64;
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es 3; Conserv
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GenCore version 5.1.6 Copyright (c) 1993 - 2004 Compugen Ltd.

OM protein - protein search, using sw model

Run on:

2, 2004, 09:56:18; Search time 46 Seconds
 (without alignments)
92.135 Million cell updates/sec April

US-10-009-049-1 84 Perfect score:

1 CRAHSFGSPRPLPVV 15 Sequence:

BLOSUM62 Gapop 10.0 , Gapext 0.5 Scoring table:

Total number of hits satisfying chosen parameters: 1586107 seqs, 282547505 residues

Minimum DB seq length: 0 Maximum DB seq length: 15

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

A Geneseq 29Jan04:* Database :

geneseqp2003as:* geneseqp2003bs:* geneseqp1980s;* geneseqp2000s:* geneseqp2001s:* geneseqp2002s:* geneseqp1990s:* geneseqp2004s:* Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

	Description	Aab59249 Peptide r	20	3251	N-termin	69 Proline-	Abq72660 Human K-r	Н	Aaw94619 Src homol	Aaw39011 Peptide r	Aab06977 Epitope #		Abm00140 Amylase n			2 SH3 synt	295 Huma	Aaw38981 Peptide r	Aam06985 CYPIB1 HL	Aam08985 HLA-A *02	Aam12950 HLA-A26 d	Aam08833 HLA-A1 de	Aam09489 HLA-B *07	m	Aab21272 ADAM 10/K	4
ST THE STORY	ID	AAB59249	AAB59250	AAB59251	ABR44350	AAW05069	ABG72660	AAW11111	AAW94619	AAW39011	AAB06977	ABM00138	ABM00140	AAW11105	AAW11098	AAW25512	AAM53295	AAW38981	AAM06985	AAM08985	AAM12950	AAM08833	AAM09489	AAU75613	AAB21272	AAP60604
	DB	4	4	4,	9	N	ιV	7	7	7	m	4	4	7	7	~	4	7	4	4	4	4	4	Ŋ	'n	н
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	Aam12240 HLA-B8 no Aam11714 HLA-A26 n		Aam11998 HLA-B *15
AAR93295 AAR9312 AAW22700 AAW2517 AAW98385 AAM98384 AAW98958 AAW38958 AAW38958 AAW38958 AAW3030 AAM11185 AAM11185 AAM10775 AAM09513	AAM12240 AAM11714	AAM12099 AAM08694	AAM11998
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0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0	4 4 2 2	44.	45

ALIGNMENTS

capable of modulating a fibrin fragment E activity. The perides and modulators are useful for inhibiting stimulation of cell proliferation induced by a fibrin degradation product. The antibody or binding fragment is also useful for identifying the active site of the fibrin fragment E receptor. The peptides are particularly useful in treating cancer, psocials restenosis, rheumatoid arthritis, diabetes, renal diseases or psoriasis. The antibodies are also useful for modulating the activity of the peptide New peptides capable of modulating a fibrin fragment E activity, useful for inhibiting uncontrolled cell proliferation, e.g. for treating cancer, vascular restenosis, atherogenesis or psoriasis, and for wound healing. The present invention relates to a peptide or its fragment, which is Fibrin fragment E; cell proliferation; cancer; vascular restenosis; arthritis; diabetes; renal disease; psoriasis. Peptide recognised by gene VIII library clone 45. Stirk CM; AABS9249 standard; peptide; 15 AA. Claim 1; Page 49; 59pp; English. 07-JUN-2000; 2000WO-GB002197. 99GB-00012994. 27-MAR-2001 (first entry) Thompson WD, (UYAB-) UNIV ABERDEEN. WPI; 2001-080581/09. WO200075175-A1. 07-JUN-1999; 14-DEC-2000. Melvin WT, Synthetic. AAB59249; RESULT 1

Sequence 15 AA;

Query Match

100.0%; Score 84; DB 4; Length 15;

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                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention relates to a peptide or its fragment, which is capable of modulating a fibrin fragment E activity. The peptides and modulators are useful for inhibiting stimulation of cell proliferation induced by a fibrin degradation product. The antibody or binding fragment is also useful for identifying the active site of the fibrin fragment E receptor. The peptides are particularly useful in treating cancer, vascular restenosis, rheumatoid arthritis, diabetes, renal diseases or psoriasis. The antibodies are also useful for modulating the activity of
                                                                                                                                                                                                                                                                                                                                                                         New peptides capable of modulating a fibrin fragment E activity, useful for inhibiting uncontrolled cell proliferation, e.g. for treating cancer, vascular restenosis, atherogenesis or psoriasis, and for wound healing.
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                                                                                                                                                                                Fibrin fragment E; cell proliferation; cancer; vascular restenosis;
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0
            0;
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           Indels
                                                                                                                                                             Peptide #2 recognised by gene VIII library clone 45.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Peptide #3 recognised by gene VIII library clone 45.
 Pred. No. 3.8e-06;
Mismatches 0;
                                                                                                                                                                                            arthritis; diabetes; renal disease; psoriasis.
                                                                                                                                                                                                                                                                                                                                     Stirk CM;
                                                                                                 AAB59250 standard; peptide; 15 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAB59251 standard; peptide; 15 AA.
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0
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100.0%;
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                               12
                                                 1 CRAHSFGSPRPLPVV 15
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                                                                                                                                         (first entry)
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 Best Local Similarity 100.
Matches 15; Conservative
                              CRAHSFGSPRPLPVV
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                                                                                                                                                                                                                                                                                                                                    Thompson WD,
                                                                                                                                                                                                                                                                                                                (UYAB-) UNIV ABERDEEN
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es 14; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 15 AA;
                                                                                                                                                                                                                                                                                            07-JUN-1999;
                                                                                                                                        27-MAR-2001
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                                                                                                                                                                                                                                                     14-DEC-2000
                                                                                                                                                                                                                                                                                                                                    Melvin WT,
                                                                                                                                                                                                              Synthetic.
                                                                                                                     AAB59250;
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The present invention relates to a peptide or its fragment, which is capable of modulating a fibrin fragment B activity. The peptides and modulators are useful for inhibiting stimulation of cell proliferation induced by a fibrin degradation product. The antibody or binding fragment is also useful for identifying the active site of the fibrin fragment receptor. The peptides are particularly useful in treating cancer, vascular restenosis, rhematoid arthritis, diabetes, renal diseases or psoriasis. The antibodies are also useful for modulating the activity of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New peptides capable of modulating a fibrin fragment B activity, useful for inhibiting uncontrolled cell proliferation, e.g. for treating cancer, vascular restenosis, atherogenesis or psoriasis, and for wound healing.
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Fibrin fragment E; cell proliferation; cancer; vascular restenosis; arthritis; diabetes; renal disease; psoriasis.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 75; DB 4; Length 10, Pred, No. 9.5e-05;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     Stirk CM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABR44350 standard; peptide; 15 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 1; Page 49; 59pp; English.
                                                                                                                                                                                                                                                                    07-JUN-2000; 2000WO-GB002197.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              89.3%;
93.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  10-APR-2001; 2001CN-00105912.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 CRAHSFVSPRPLPVV 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 CRAHSFGSPRPLPVV 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          14-JUL-2003 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Thompson WD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                       (UYAB-) UNIV ABERDEEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2001-080581/09.
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                                                                                                                                              WO200075175-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 15 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mao Y, Xie Y;
                                                                                                                                                                                                                                                                                                                               07-TUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CN1380320-A.
                                                                                                                                                                                                         14-DEC-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              the peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20-NOV-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                               Melvin WT,
                                                                                             Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
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                                                                                                                The present invention discloses a polypeptide-guanosine triphosphatase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New DNA encoding proteins that bind to Src-homology domain 3 - used to inhibit signal transduction, e.g. in cases of T cell activation.
                                                                                                                                  for
                                                                                                                                                                                                                                                                            Gaps
                                                                                                                               activator protein-10.01. The invention also discloses the method for curing several disceases, such as squamobasal cell carcinoma of skin, osteosarcoma, leukemia and teratoma by using said polypeptide. The present sequence represents the N-terminal sequence of guanosine
                                          polypeptide-guanosine triphosphatase activator protein -10.01 and symucleotide for coding this polypeptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       7..13
/label= SH3_binding_motif
/note= "corresponds to consensus sequence RPLPXXP"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Proline-rich SH3 binding peptide fuseSB-4.51 from random library.
                                                                                                                                                                                                                                                                            0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Jaye MC;
                                                                                                                                                                                                                                                6; Length 15;
                                                                                                                                                                                                                                                                            2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Src homology domain; SH3 domain; oncogene; consensus; random display library.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             South VJ,
                                                                                                                                                                                                                                                DB
                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                              Score 38;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cheadle C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (RHON ) RHONE POULENC RORER PHARM INC
                                                                                                                                                                                       triphosphate activator protein 10.01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers
                                                                                    Example 6; Page 21; 33pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                                                                      AAW05069 standard; peptide; 15 AA.
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                                                                                                                                                                                                                                              45.2%;
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Searfoss GH;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                       polynucleotide for coding
                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                      4 HSFGSPRPLP 13
                                                                                                                                                                                                                                                                                                                                 4 HSLGSLQPLP 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                (revised)
                                                                                                                                                                                                                               Query Match
Best Local Similarity
77, Conserve
               WPI; 2003-222552/22
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                                                                                                                                                                                                                   Sequence 15 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               25-MAR-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          French SM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                  AAW05069;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention discloses a new kind of polypeptide, human K-ras proto-oncogene protein 10.67, polynucleotides encoding the polypeptide and a DNA recombination process to produce the polypeptide. The present invention also discloses applying the polypeptide in treating various diseases, such as malignant tumours, haemopathy, development disorder, human immunodeficiency virus (HTV) infection, immunological diseases and various inflammations. The present invention also discloses the antagonist resisting the polypeptide and its treatment effect. The present invention also discloses application of the polynucleotides encoding human K-ras proto-oncogene protein 10.67, The present sequence represents human K-ras proto-oncogene protein 10.67, N-terminus, used in an ELISA (enzyme-linked immunosorbent assay) experiment
                    18
identified within a protein isolated by screening a lambda-lox mouse embryo CDNA library with a Src SH3 domain probe. The present sequence is that of one of the positive peptides identified from the fuse 5B random display library. (Updated on 25-MAR-2003 to correct PI field.)
                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human, K-ras 10.67; proto-oncogene; tumour; haemopathy;
development disorder; human immunodeficiency virus infection; HIV;
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                                                                                                                             Length 15;
                                                                                                                                                              Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                          Human K-ras 10.67 proto-oncogene protein, N-terminus.
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ilarity 85.7%; Pred. No. 1.1e+02;
Conservative 1; Mismatches 0;
                                                                                                                           Score 37; DB 2;
Pred. No. 74;
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (BODE-) BODE GENE DEV CO LTD SHANGHAI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  immunological disease; inflammation.
                                                                                                                                                                                                                                                                                                                     ABG72660 standard; peptide; 15 AA.
                                                                                                                         44.0%;
70.0%;
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                                                                                                       Query Match
Best Local Similarity 70.0
                                                                                                                                                                                             4 HSFGSPRPLP 13
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Best Local Similarity
6; Conserve
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2002-658691/71.
                                                                                                                                                                                                                                1 HSHFHPRPLP
                                                                                      Sequence 15 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 15 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mao Y, Xie Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ното варіепв
                                                                                                                                                                                                                                                                                                                                                                                         26-FEB-2003
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7 GSPRPLP

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WO9730074-A1
                                         Synthetic
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                  AAWI1098-WI1124 are peptides that bind to the Src SH3 domain. The SH3 binding peptides are useful in modulating signal transduction pathways at the cellular level (especially protein tyrosine kinase-mediated), modulating oncogenic protein activity, or providing compounds for the development of drugs with the ability to modulate broad classes, as well as specific classes, of proteins involved in signal transduction and also for regulating the processing, trafficking or translation of RNA. Conjugates of the peptides with detectable labels or imaging agents are useful for imaging cells, tissues and organs in which Src or Src-related
                                                                                                                                                                                                                                                                                                                                                      Peptide with binding affinity for Src homology region 3 (SH3) domains of proteins - useful for e.g. modulating signal transduction pathways at the cellular level, esp. protein tyrosine kinase-mediated.
                                                                                                              Src SH3 domain-binding peptide used in signal transduction modulation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Src homology 3 domain; SH3 domain; peptidomimetic; mimetic; modulator;
                                                                                                                                Src; SH3; Src homology region 3; binding affinity; oncogenic protein;
protein tyrosine kinase; signal transduction; RNA processing;
trafficking; translation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       41.7%; Score 35; DB 2; Length 13; 62.5%; Pred. No. 1.38+02; Live 3; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                S.
                                                                                                                                                                                                                                                                                                                 Der
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Src homology 3 domain binding nonamer peptide #1.
                                                                                                                                                                                                                                                                                                                Quilliam LA,
                                                 AAW11111 standard; peptide; 13 AA.
                                                                                                                                                                                                                                                                                                                                                                                                 Claim 38; Page 87; 116pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAW94619 standard; peptide; 9 AA.
                                                                                                                                                                                                                                                                                                                Thorn JM,
                                                                                                                                                                                                                                                            94US-00278865.
95US-00483555.
                                                                                                                                                                                                                                                                                           (UYNC-) UNIV NORTH CAROLINA.
                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              5; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  proteins are expressed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   8 SPRPLPVV 15
                                                                                                                                                                                                                                                                                                                                   WPI; 1996-117151/12.
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                                                                                                                                                                                                                                                                                                                Kay BK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
GSPOPLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 13 AA;
                                                                                                                                                                                                WO9603649-A1.
                                                                                                                                                                                                                                                                       07-JUN-1995;
                                                                                          25-JUN-1997
                                                                                                                                                                                                                   08-FEB-1996
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                                                                                                                                                                                                                                                                                                               Sparks AB,
                                                                                                                                                                           Synthetic.
                                                                     AAW11111;
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                                      AAW11111
                           RESULT
                                                           W X X B X B X X X X X
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The present invention describes spiro[Pyrrolidine-2':3-piperidin-2-ones], their fused ring analogues, and salts of both, all as individual compounds or their libraries. They are peptidominetics, and interact with protein Src homology 3 (SH3) Gomains, involved in protein-protein interactions in signal transduction. They cause disruption of the pathway in particular, they inhibit the protein tyrosine kinase system involved in cell growth regulation. The cells may be microbial, e.g. pathogenic bacterial or fungal cells responsible for infections, or of transformed cells, e.g. responsible for neoplastic conditions. Specific examples of Src proteins with SH3 domains are those known by the abbreviations p85, Fyn, Lyn, Hck, Syk, Grb2, Gap, STAT, p47-phox, p67-phox, and Brk. The present sequence represents a nonamer peptide used in an example from the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    spiro[Pyrrolidine-2':3-piperidin-2-ones] and their fused ring analogues used, e.g. to modulate signal transduction protein interactions, inhibit cell growth and for use in treating infections and cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cortactin; SH3 domain; binding peptide; Src homology region 3; tyrosine kinase; immune response; lymphokine; interleukin 1; Nck; Abl; PLCgamma; p53bp2; Crk; Yes; Grb2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ..
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beta-turn template; spiro[Pyrrolidine-2':3-piperidin-2-ones]; fused ring analogue; signal transduction protein interaction; inhibit cell growth; infection; cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                40.5%; Score 34; DB 2; Length 9; 100.0%; Pred. No. 1.4e+06; Live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                  Location/Qualifiers
                                                                                                                                                                                                                                                                                                          /note= "acetylated"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (CADU-) CADUS PHARM CORP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1999-080804/07.
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                                                                                                                                                                                                                                                                  Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        28-MAY-1998;
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25-NOV-1998;
            04-OCT-1999;
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                                                          Roggen EL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Roggen EL,
                                                                                                                                                                                                                                                                                                                   (AAB06998)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABM00138;
                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                          Matches
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                                                                                                                                                                                                                                                 The present sequence represents a peptide which resembles a Src homology region 3 (SH3) binding peptide. SH3 binding peptides are selected from:

(a) peptides which bind the SH3 domain of Cortactin; (b) peptides which bind the Mck; (c) peptides which bind the SH3 domain of Nck; (c) peptides which bind the SH3 domain of Ab1; (d) peptides which bind the SH3 domain of PLC gamma; (f) peptides which bind the SH3 domain of peptides which bind the SH3 domain of Crk; (h) peptides which bind the SH3 domain of Crk; (h) peptides which bind the SH3 domain of Crk; (h) peptides which bind the SH3 domain of Grb; reputides which bind the amino-terminal SH3 domain of Grb. The purified binding peptides can be used in the method to identify inhibitors of their binding to their respective SH3 domains, which could be used to modulate the pharmacological activity of proteins or polypeptide containing the SH3 domain. The peptides can also be used to activate Src or Src-related
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Low allergenic protein variant; reduced immunogenicity; allergy prevention; epitope; personal care product; enzyme; food industry; textile industry; pharmaceutical; JE-1; natalase; amylase.
                                                                                                                                                                                                                                                                                                                                                                                                            protein tyrosine kinases, to stimulate the immune response by increasing the production of certain lymphokines, e.g. tumour necrosis factor-alpha and interleukin-1, or to deliver a conjugated molecule to certain cellular compartments containing src or Src related proteins
                                                                                                                                                                            Src homology region 3 binding peptide - used to activate Src tyrosine kinase(s) and to stimulate immune response by increasing production of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                      Fowlkes DM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ..
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 40.5%; Score 34; DB 2; Length 15; 100.0%; Pred. No. 2.2e+02; ive 0; Mismatches 0; Indels
                                                                                                                    Der CJ,
                                                                                                                   Quilliam LA,
                                                                                                                                                                                                      certain lymphokine(s), e.g. interleukin-1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Epitope # 7 from JE-1 (Natalase/ amylase).
                                                                                                                                                                                                                           Claim 22; Page 93; 131pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAB06977 standard; peptide; 11 AA.
                                                                                                                   Thorn JM,
                                  97WO-US002298
                                                         96US-00602999
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                                                                                           UYNC-) UNIV NORTH CAROLINA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          6; Conservative
                                                                                (CYTO-) CYTOGEN CORP
                                                                                                                   Sparks AB, Kay BK,
                                                                                                                                                       WPI; 1997-424972/39.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                9 PRPLPV 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRPLPV 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 15 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO200026230-A1
                                 14-FEB-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Unidentified.
                                                       16-FEB-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 30-OCT-1998;
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          21-AUG-1997
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                                                                                                                               Rider JE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAB06977;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
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The present invention relates to selecting a protein variant having reduced immunogenicity as compared to a parent protein. The selected proteins can be enzymes or peptide hormones. The proteins which are selected are low allergenic proteins which can be used to prevent cases of allergy in susceptible individuals. They can be used in the food or textile industries, housekeeping, medicine, e.g. proteins used in epitope from JE-1 (Natalase/ amylase) parent protein. This sequence as derived by screening a random peptide display package library with antibodies raised against JE-1. This sequence was then subjected to genetical engineering mutations to reduce its immunogenicity. This sequence was used in a sequence alignment to determine an epitope pattern
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                                                                                                                                                                                                                                                                                                                   Novel method for selecting low allergenic protein variants which are useful in industrial, food, and pharmaceutical preparations.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               39.3%; Score 33; DB 3; Length 11; 60.0%; Pred. No. 2.38+02; ive 1; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Amylase natalase antibody binding peptide sequence 24.
                                                                                                                                                                                                                                                                                                                                                                                                                            Example 2; Page 66; 91pp; English.
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98DK-00001645.
                                 99DK-00001417.
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10-MAY-2000; 2000US-0203345P.
28-FEB-2001; 2001DK-00000327.
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                                                                                                     (NOVO ) NOVO NORDISK AS
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                                                                                                                                                                        Olsen AA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Ernst S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (NOVO ) NOVOZYMES AS
                                                                                                                                                                                                                                                WPI; 2000-365562/31.
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The invention relates to selecting a protein variant having modified immunogenicity, compared to a parent protein, comprising using the antibody binding sequence to localise epitope sequences on the three dimensional structure of the parent protein and defining an epitope area including amino acids within 5 Angstrom of the epitope amino acids. The method is useful for identifying structural epitopes on the 3-dimensional surface of commercial and environmental allergens. Compositions containing the protein variants are used as vaccines, detergents and personal care compositions, e.g. shampoo, balsam, hair conditioners, hair waring compositions, hair dyeing compositions, hair dyeing compositions, hair dyeing compositions, shir cream, sunscreen, shaving compositions, hair spray, chewing gum, skin cream, sunscreen, shaving foam, cream soap, skin milk or foundation. The present sequence is that of an antibody binding peptide sequence related to the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Allergen; protein coordinate data; vaccine; antiallergic; immunogenicity;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to selecting a protein variant having modified immunogenicity, compared to a parent protein, comprising using the antibody binding sequence to localise epitope sequences on the three dimensional structure of the parent protein and defining an epitope area
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           produce vaccines, detergents and personal care compositions, involves localizing epitope sequences on the three-dimensional structure of a
                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
 epitope sequences on the three-dimensional structure of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Selecting protein variants having modified immunogenicity, used to
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                                                                                                                                                                                                                                                                                                                                                                                           39.3%; Score 33; DB 4; Length 12; 60.0%; Pred. No. 2.5e+02; tive 1; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Amylase natalase antibody binding peptide sequence 26.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          detergent; personal care composition; cosmetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 1; Page 148; 513pp; English.
                                                     Example 1; Page 148; 513pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2000US-0203345P.
2001DK-00000327.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ernst S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2001-626552/72.
                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
                                                                                                                                                                                                                                                                                                                                                       Sequence 12 AA;
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Matches
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including amino acids within 5 Angstrom of the epitope amino acids. The method is useful for identifying structural epitopes on the 3-dimensional surface of commercial and environmental allergens. Compositions containing the protein variante are used as vaccines, detergents and personal care compositions, e.g. shampoo, balsam, hair conditioners, hair waving compositions, hair dyeing compositions, hair tonic, hair liquid, hair cream, hair rinse, hair spray, chewing gum, skin cream, sunscreen, shaving foam, cream soap, skin milk or foundation. The present sequence is that of an antibody binding peptide sequence related to the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Peptide with binding affinity for Src homology region 3 (SH3) domains of proteins - useful for e.g. modulating signal transduction pathways at the cellular level, esp. protein tyrosine kinase-mediated.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            modulating oncogenic protein activity, or providing compounds for the development of drugs with the ability to modulate broad classes, as well as specific classes, of proteins involved in signal transduction and also for regulating the processing, trafficking or translation of RNA. Conjugates of the peptides with detectable labels or imaging agents are useful for imaging cells, tissues and organs in which Src or Src-related
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          binding peptides are useful in modulating signal transduction pathways at
the cellular level (especially protein tyrosine kinase-mediated),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Src SH3 domain-binding peptide used in signal transduction modulation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Src; SH3; Src homology region 3; binding affinity; oncogenic protein; protein tyrosine kinase; signal transduction; RNA processing;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAW11098-W11124 are peptides that bind to the Src SH3 domain. The SH3
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Pred. No. 2.5e+02;
1; Mismatches 3; Indels
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Pred. No. 2.7e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                           AAW11105 standard; peptide; 13 AA.
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                                                                                                                                                                                                                      39.3%;
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95US-00483555.
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62.5%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1996-117151/12.
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Best Local Similarity
                                                                                                                                                                                                                                         Best Local Similarity
                                                                                                                                                                                     Sequence 12 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  25-JUN-1997
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               of
the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Peptide with binding affinity for Src homology region 3 (SH3) domains proteins - useful for e.g. modulating signal transduction pathways at cellular level, esp. protein tyrosine kinase-mediated.
                                                                                                                                                                                                                                Src SH3 domain-binding peptide used in signal transduction modulation.
 Gaps
                                                                                                                                                                                                                                                             Src, SH3; Src homology region 3; binding affinity; oncogenic protein; protein tyrosine kinase; signal transduction; RNA processing;
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 Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Quilliam LA,
 3; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAW25512 standard; peptide; 14 AA.
                                                                                                                                     AAW11098 standard; peptide; 14 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 32; Page 79; 116pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Thorn JM,
                                                                                                                                                                                                                                                                                                                                                                                                                      95WO-US009382.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   94US-00278865,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  95US-00483555.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (UYNC-) UNIV NORTH CAROLINA.
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                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                               trafficking; translation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity 62.5
les 5; Conservative
 5; Conservative
                             8 SPRPLPVV 15
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                                                        2 TPRPLPML 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sparks AB, Kay BK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 14 AA;
                                                                                                                                                                                                                                                                                                                                                        WO9603649-A1
                                                                                                                                                                                                                                                                                                                                                                                                                    24-JUL-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 22-JUL-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   07-JUN-1995;
                                                                                                                                                                                                 25-JUN-1997
                                                                                                                                                                                                                                                                                                                                                                                     08-FEB-1996.
                                                                                                                                                                                                                                                                                                                           Synthetic
                                                                                                                                                                    AAW11098;
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                                                                                                      RESULT 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              protein tyrosine kinases, to stimulate the immune response by increasing the production of certain lymphokines, e.g. tumour necrosis factor-alpha and interleukin-1, or to deliver a conjugated molecule to certain cellular compartments containing Src or Src related proteins
                                                Cortactin; SH3 domain; binding peptide; Src homology region 3; tyrosine kinase; immune response; lymphokine; interleukin 1; NCk; Abl; PLCgamma; p53bp2; Crk; Yes; Grb2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Src homology region 3 binding peptide - used to activate Src tyrosine kinase(s) and to stimulate immune response by increasing production of certain lymphokine(s), e.g. interleukin-1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         pharmacological activity of proteins or polypeptide containing the SH3 domain. The peptides can also be used to activate Src or Src-related
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                            Fowlkes DM;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 14;
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62.5%; Pred. No. 2.9e+02;
trive 3; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                          Quilliam LA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Search completed: April 2, 2004, 10:08:41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Fig 7; 131pp; English.
                                                                                                                                                                                                                                                                                                                                                                                          Sparks AB, Kay BK, Thorn JM,
              SH3 synthetic binding peptide.
                                                                                                                                                                                                                                                  97WO-US002298.
                                                                                                                                                                                                                                                                                        96US-00602999
                                                                                                                                                                                                                                                                                                                             (CYTO-) CYTOGEN CORP.
(UYNC-) UNIV NORTH CAROLINA.
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Matches 5; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 14 AA;
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                                                                                                                                                                                                                                                  14-FEB-1997;
                                                                                                                                                                                                                                                                                        .6-FEB-1996;
                                                                                                                                                                      WO9730074-A1
                                                                                                                                                                                                             21-AUG-1997.
                                                                                                                                  Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                            Rider JE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
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GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
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OM protein - protein search, using sw model

Run on:

April 2, 2004, 10:05:03 ; Search time 11 Seconds (without alignments) 131.170 Million cell updates/sec

US-10-009-049-1 84 1 CRAHSFGSPRPLPVV 15

Title: Perfect score:

Scoring table: Sequence:

283366 seqs, 96191526 residues Searched:

BLOSUM62 Gapop 10.0 , Gapext 0.5

2522 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 15

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

78:* PIR Database

1: pirl:* 2: pir2:* 3: pir3:* 4: pir4:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

	ption	hypothetical prote	sin-conve	protein OF200053 -	iq H chain V-D-J r	in-potent	bradykinin-potenti	kinesin light chai	photosystem II oxy	T-cell receptor be			_	neuropeptide calla		photosystem II oxy	tidy	Ig heavy chain CRD	ythrocy		locustamyotropin I	platelet-derived q	dermorphin - Rohde	orf 1 rara 5'-regi	triacyldlycerol li	173K exoantigen -		ptor	l receptor b	ransferase
SUMMARIES		S27140	XAVI9B	PA0089	PH1618	$\boldsymbol{\sigma}$	D37196	S43170	PA0013	PH1457	1O	PS0452	S	D47393	PL0157	G44957	I84603	PT0238	XGHUE	JH0460	B61620	A22789	A61324	S15594	PX0030	I33098	S01122	PH1459	PH1468	12
	DB	5	Н	~	7	7	N	~	7	C)	N	7	7	7	Ŋ	ď	N	~	~	N	N	N	Н	4	N	~	~	7	~	4
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de	Query	32.1	ä	ä	28.6	8	8	θ.	8	27.4	27.4	ů.	è.	26.2	ġ.	9	ß	S	25.0	S	rU.	S	3	\sim	m	3	m	m	'n	m
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formate dehydrogen	orf19 3' of eryK -	hypothetical prote	mixed lymphocyte r	leucokinin VI - Ma	xylose isomerase (angiotensin-conver	bradykinin-potenti	substance P - chic	substance P - rain	substance R - Atla	probable antigen 5	probable substance	T-cell receptor be	T-cell receptor be	T-cell receptor de
S59492	A40634	B32800	PX0031	JS0316	A31576	XAV16B	XASNBA	JN0023	S23308	S23 306	E41476	833300	\$26559	S26554	B49033
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23.8	23.8	23.8	23.8	22.6	22.6	22.6	22.6	22.6	22.6	22.6	22.6	22.6	22.6	22.6	22.6
20	20	20	20	19	19	19	19	19	19	19	19	19	19	19	19
30	31	32	33	34	35	36	3.7	38	39	40	41	42	43	44	45

ALIGNMENTS

	hypothetical protein 1 estrogen receptor 5'-region - human C.Species: Homo sapiens (man) C.Date: OS-Mar-1994 #sequence_revision 12-May-1995 #text_change 18-Aug-2000 C.Accession: S27140	R.Keaveney, M.; Klug, J.; Gannon, F. DNA Seq. 2, 347-358, 1992 A;Title: Sequence analysis of the 5' flanking region of the human estrogen receptor gene A;Akeference number: S27140; MUID:93075998; PMID:1476547 A;Accession: S27140	A.Status: translation not shown A.Molecule type: DNA A.Mesdüuss: 1-14 «KEA» A.Gross-references: EMBL:X62462; NID:g31201; PIDN:CAA44319.1; PID:g31202 C.Superfamily: unassigned leader peptides	<pre>Query Match 32.1%; Score 27; DB 2; Length 14; Best Local Similarity 100.0%; Pred. No. 3.5e+02; Matches 5; Conservative 0; Mismatches 0; Indels 0; Gaps 0; 2 RAHSF 6 2 RAHSF 6</pre>	
RESULT 1 S27140	hypothetica C;Species: C;Date: 05-	R.Keaveney, M.; Klus DNA Seq. 2, 347-358 A.Title: Sequence an A.Keference number: A.Accession: S27140	A;Status: t A;Molecule A;Residues: A;Cross-ref C;Superfami	Query Mat. Best Loca Matches Qy	

angiotensin-converting enzyme inhibitor V-9 - jararaca C;Species: Bothrops jararaca (jararaca) C;Date: 13-Jul-1981 #sequence_revision 13-Jul-1981 #text_change 08-Dec-1995

Cyaccession: A01253
RyOndetti, M.A.; Williams, N.J.; Sabo, E.F.; Pluscec, J.; Weaver, E.R.; Kocy, O. Biochemistry 10, 4033-4039, 1971
Biochemistry 10, 4033-4039, 1971
Biochemistry 10, 4033-4039, 1971
A;Reference number: A90356; MUD:72118526; PMID:4334402
A;Recession: A01253
A;Rocession: A01253
A;Rocession: A01253
A;Rocession: A01253
C;Comment: This peptide also potentiates bradykinin by inhibiting the kinases that inact C;Comment: This peptide also potentiating peptide
C;Comment: This peptide also potentiating peptide
C;Superfamily: bradykinin-potentiating peptide
C;Keywords: angiotensin-converting enzyme inhibitor; pyroglutamic acid
F;1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental

Gaps . 0 Query Match 31.0%; Score 26; DB 1; Length 13; Best Local Similarity 55.6%; Pred. No. 4.8e+02; Matches 5; Conservative 1; Mismatches 3; Indels

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7 GSPRPLPVV 15 3 GWPRPGPEI 11 ð g

RESULT 3

31.0%; Score 26; DB 2; Length 13; 80.0%; Pred. No. 4.8e+02; tive 1; Mismatches 0; Indels

Local Similarity 80.0 les 4; Conservative

Matches

Query Match

7 GSPRP 11

5 GAPRP 9

RESULT 4

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C;Species: Home sapiens (man)
C;Species: Home sapiens (man)
C;Date: 13-Jan-1995 #sequence_revision 18-Aug-2000 #text_change 18-Aug-2000
C;Accession: S43170
R;Chernajovsky, Y.; Brown, A.; Jones, T.A.; Sheer, D.
Submitted to the EMBL Data Library, December 1992
A;Description: Promoter first exon/intron characterization and chromosomal location of the A;Reference number: S43170
A;Reference number: S43170
A;Reference Letype: Dyanary
A;Residues: 1-12 CCHE>
                                                                                                                                                                                                                                                                                                                                                             C'Accession: D37196
R'Cintra, A.C.O.; Vieira, C.A.; Giglio, J.R.
J. Protesin Chem. 9, 221-227, 1990
A;Title: Primary structure and biological activity of bradykinin potentiating peptides for A;Reference number: A37196; MuID:90351557; PMID:2386615
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 C;Accession: PA0013
R;Kamo, M.; Kawakami, T.; Miyatake, N.; Tsugita, A.
Submitted to JTPID, July 1994
A;Description: Separation and characterization of Arabidopsis proteins by two-dimensiona.
A;Reference number: PA0001
A;Accession: PA0013
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   photosystem II oxygen-evolving complex protein 2 - Arabidopsis thaliana (fragment) C;Species: Arabidopsis thaliana (mouse-ear cress) C;Date: 30-Jun-1992 #sequence_revision 06-Jan-1995 #text_change 23-Mar-1995
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                                                                                                                                                                                                                          bradykinin-potentiating peptide 4 - island jararaca
C;Species: Bothrops insularis (island jararaca)
C;Date: 14-Feb_1992 #sequence_revision 01-Dec-1992 #text_change 05-Aug-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A;Status: preliminary
A;Molecule type: protein
A;Residues: 1-11 <CIN>
C;Keywords: pyroglutamic acid
F;1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 28.6%; Score 24; DB 2; Length 14; Best Local Similarity 50.0%; Pred. No. 1.1e+03; Matches 4; Conservative 2; Mismatches 2; Indels
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A;Residues: 1-14 «KAM»
A;Expeziduental source: leaf
C;Keywords: photosynthesis; photosystem II
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               kinesin light chain - human (fragment)
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Best Local Similarity
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GPPRP 7
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                                                                                                                                                               protein QF200053 - fungus (Fusarium sporotrichioides) (fragment)
C;Species: Pusarium sporotrichioides
C;Date: 20-Peb-1995 #sequence_revision 20-Feb-1995 #text_change 23-Mar-2001
C;Accession: PA0009
E;Chow, L.P.; Fukaya, N.; Sugiura, Y.; Ueno, Y.; Tabuchi, K.; Tsugita, A.
Submitted to JIPID, October 1994
A;Reference number: PA0051
A;Reference number: PA0018
A;Accession: PA0089
A;Accession: PA0089
A;Residues: 1-13 <CHO>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CACCESSION: PHIGS
Ribevinson, D.A.; Campos-Torres, J.; Leder, P.
J. Exp. May 317-329, 1993
A; Title: Molecular characterization of transgene-induced immunodeficiency in B-less mice
A; Reference number: PHIS80: MUID:93301609; PMID:8315387
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        bradykinin-potentiating peptide 3 - island jararaca (Species: Bothrops insularis (island jararaca) (Species: Bothrops insularis (island jararaca) (Species: 14-Pb-1932 #sequence_revision 01-Dec-1992 #text_change 05-Aug-1994 (SAccession: C37196 (SAccession: C37196 (SAccession: C37196 (SAccession: C37196 (SACCESSION: C37196 (SACCESSION: CASTILIA) (SACCESSION: SACCESSION: SACCESS
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C;Species: Mus musculus (house mouse)
C;Date: 02-Jun-1994 #sequence_revision 02-Jun-1994 #text_change 17-Mar-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A,Status: preliminary
A;Molecule type: protein
A;Molecule type: protein
A;Residues: 1-11 <CIN>
C;Keywords: pyroglutamic acid
F;1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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, 0

Query Match 28.6%; Score 24; DB 2; Length 8; Best Local Similarity 42.9%; Pred. No. 2.8e+05; Matches 3; Conservative 1; Mismatches 3; Indels

A, Residues: 1-8 <LEV>
A, Experimental source: bone marrow pre-B lymphocyte C; Keywords: immunoglobulin

A; Molecule type: DNA

.; 0

28.6%; Score 24; DB 2; Length 11; 80.0%; Pred. No. 8.5e+02; tive 0; Mismatches 1; Indels

Query Match
Best Local Similarity 80.0°
Matches 4; Conservative

A; Accession: C37196

7 GSPRP 11

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dermorphin (Lys-7) (validated) - two-colored leaf frog
C;Species: Phyllomedusa bicolor (two-colored leaf frog)
C;Date: 10-Dec-1993 #sequence_revision 13-Mar-1997 #text_change 18-Aug-2000
C;Accession: 53662
R;Mignogna, G;; Severini, C;; Simmaco, M.; Negri, L.; Falconieri Erspamer, G;; Kreil, G.
FEBS Lett. 302, 151-154, 1992
A;Title: identification and characterization of two dermorphins from skin extracts of th
A;Reference number: S21152; MUID:92339502; PMID:1633846
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             C;Accession: D47393 T. Scott. A.G.; Yu, C.G.; Yagi, K.J.; Tobe, S.S.; Thorpe, A. Broc. Natl. Acad. Sci. U.S.A. 90, 2456-2460, 1993 A;Title: Callatostatins: neuropeptides from the blowfly Calliphora vomitoria with sequen A;Reference number: A47393; MUID:93211980; PMID:8460157
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ig kappa chain V-II region (anti-myelin-associated glycoprotein, PEC) - human (fragment)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A,Status: preliminary
A,Molecule type: protein
A,Residues: 1-7 <MIG>
C,Superfamily: dermorphin precursor; dermorphin precursor amino-terminal homology
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              .
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C;Species: Calliphora vomitoria
C;Date: 16-Feb-1994 #sequence_revision 18-Nov-1994 #text_change 28-Apr-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 C;Species: Homo sapiens (man)
C;Date: 20-Feb-1995 #sequence_revision 20-Feb-1995 #text_change 07-Feb-1997
C;Accession: PL0157; C61458
                                                                                                                                                                                                                                                                 Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          26.2%; Score 22; DB 2; Length 8; 66.7%; Pred No. 2.8e+05;
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A;Note: sequence extracted from NCBI backbone (NCBIP:128479)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1; Mismatches
                                               A,Molecule type: protein
A,Residues: 1-15 <TSU>
A,Experimental source: bran, strain Nihonbare
C,Comment: molecular weight 32K, pI 5.3.
                                                                                                                                                                       Query Match
Best Local Similarity 45.5%,
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                                                                                                                                                                                                                                                                                                                    5 SFGSPRPLPVV 15
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A; Reference number: PS0208
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A; Molecule type: protein A; Residues: 1-8 < DUV>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             4 HSFGSPR 10
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                             A; Accession: PS0452
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                                                                                                                                                                                                                                               C;Species: Mus musculue (house mouse)
C;Date: 10-Mar-1994 #sequence_revision 10-Mar-1994 #text_change 17-Mar-1999
C;Accession: PH1457; S26542
R;Casanova, J.L., Martinon, F.; Gournier, H.; Barra, C.; Pannetier, C.; Regnault, A.; Kc J. Exp. Med. 177, 811-820, 1993
A;Title: T cell receptor selection by and recognition of two class I major histocompatible. A;Reference number: PH1430; MUID:93171821; PMID:8436911
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A; Molecule type: mRNA
A; Residues: 1-12 cCAS>
A; Residues: 1-12 cCAS>
A; Residues: 1-12 cCAS>
A; Respectimental source: cytolytic T-lymphocyte, clone A3/IC1
R; Casanova, J.L.; Cerottini, J.C.; Matthes, M.; Necker, A.; Gournier, H.; Barra, C.; Wid
B, Exp. Med. 176, 439-47, 1992.
A; Title: H-2-restricted cytolytic T lymphocytes specific for HLA display T cell receptor
A; Reference number: S26512; MUID: 92364546; PMID: 1380061
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            C; Species: Fusarium sporotrichioides
C; Species: Co-Feb-1995 #sequence_revision 20-Feb-1995 #text_change 23-Mar-2001
C; Accession: PA0060
R; Chow, L.P.; Fukaya, N.; Sugiura, Y.; Ueno, Y.; Tabuchi, K.; Tsugita, A.
Bubmitted to JTPLD, October 1994
A; Description: Two dimensional polyacrylamide gel electropheresis of Fusarium sporotrich
A; Reference number: PA0051
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ó
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C;Date: 03-Feb-1994 #sequence_revision 03-Feb-1994 #text_change 11-Apr-1995
C;Accession: PS0452
Sibragita, A.; Miyatake, N.
submitted to JIPID, April 1993
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                27.4%; Score 23; DB 2; Length 12; 83.3%; Pred. No. 1.3e+03; ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                27.4%; Score 23; DB 2; Length 15; 100.0%; Pred. No. 1.7e+03; bred. No. 1.7e+03; tive. 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; Experimental source: cytolytic T-lymphocyte, clone Cw3/Cas2 C; Superfamily: immunoglobulin homology
                                                                                                                                                                                                                          T-cell receptor beta chain (clone A3/IC1) - mouse (fragment)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 32K protein 3306 - rice (strain Nihonbare) (fragment)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            5; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          C; Keywords: receptor; T-cell
                       3 AHSFGSPR 10
                                                                             6 ANVFGKPK 13
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A;Molecule type: protein
A;Residues: 1-15 <CHO>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A; Molecule type: mRNA
A; Residues: 1-12 < CA2>
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Matches
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PS0452

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photosystem II oxygen-evolving complex protein 2 - common tobacco (cv. Samsun NN) (fragm C;Species: Nicotiana tabacum (common tobacco)
C;Species: Nicotiana tabacum (common tobacco)
C;Accession: 644557
R;Takeda, S; Sato, F; Ida, K.; Yamada, Y.
R;Takeda, S.; Sato, F; Ida, K.; Yamada, Y.
R;Reference number: A44957
A;Reference number: A44957
A;Reference number: A44957
A;Reference number: A44957
A;Residues: 1-14 cTAK>
                                                                                                                                                                                                                                                               A,Accession: C61458
A;Status: preliminary
Molecule type: protein
A;Residues: 1-13 < BR2>
C;Comment: This protein is one of monoclonal IgM reactive with myelin-associated glycopz
C;Keywords: glycoprotein; heterotetramer; immunoglobulin
R;Brouet, J.C.; Dellagi, K.; Gendron, M.C.; Chevalier, A.; Schmitt, C.; Mihaesco, E. J. Exp. Med. 170, 1551-1558, 1989
A;Title: Expression of a public idiotype by human monoclonal IgM directed to myelin-assot A;Reference number: A61458; MUD:90039128; PMID:2478651
A;Accession: PL0157
A;Molecule type: protein
A;Residues: 1-13 <BRO>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match

26.2%; Score 22; DB 2; Length 13;
Best Local Similarity 71.4%; Pred, No. 2.1e+03;
Matches 5; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     26.2%; Score 22; DB 2; Length 14; 57.1%; Pred. No. 2.38+03; tive 1; Mismatches 2; Indels
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Best Local Similarity 57.1
Matches 4; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       8 SPRPLPV 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             7 SPUTLPV 13
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6 ANVFGKP 12
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GenCore version 5.1.6
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OM protein - protein search, using sw model

April 2, 2004, 10:04:38 ; Search time 31.3333 Seconds (without alignments) 151.046 Million cell updates/sec Run on:

US-10-009-049-1 Perfect score:

84 1 CRAHSFGSPRPLPVV 15 Sequence:

BLOSUM62 Gapop 10.0 , Gapext 0.5 Scoring table:

1017041 segs, 315518202 residues Searched:

3954 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 15

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database :

sp_archea:*
sp_bacteria:*
sp_bacteria:*
sp_human:*
sp_human:*
sp_invertebrate:*
sp_mammal:* SPTREMBL 25:*

sp_rodent:*
sp_virus:*
sp_vertebrate:*
sp_unclassified:* sp_rvirus:*
sp_bacteriap:*
sp_archeap:* sp_organelle:* sp_phage:* sp plant: * 101. 111. 112. 113. 114. 115.

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

STAMMARTES

			Description	•	CARROL CLADINECOGI	obcias musculu	ນໄທລະເໜ ອະເລອດ	O98696 hordenmil	095771 ofenogames				CAP//O CACTURE LIC	Ol3865 homo sapien	O9twv0 anthopleura	O8cq39 rattus norv	OBiv78 October		00,000000000000000000000000000000000000	Constant of the contract	Q9euz3 escherichia	018890 ateles belz	
SOUTHWANTES			ID		てつにまんび	QBCJA8	OBCJA9	969860	095771	095952	095879	001110	01100	Q13865	Q9TWV0	080639	08JV78	088176	OBGTGS		Q9EUZ3	018890	
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	₩	Query	Match	27.0		. 31.0	31.0	29.8	29.8	29.8	29.8	0.0		28.6	27.4	27.4	27.4	27.4	26.2		7.07	26.2	
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Q993c0 calliphora	ဌ	Q9s905 glycine max	Q8mje4 aotus azara	Q9tnq8 homo sapien	Q9qvi3 cavia (quin	homo	homo	ошо	Q9qwt0 mus musculu	Q7x566 thermus the		Osmips callimico q	O8mjp7 callithrix		O8mje3 saimiri sci			Q9qvk4 rattus sp.	rattus	Q9uj52 homo sapien	O95179 homo sapien	Q26323 lymnaea sta	P82439 nicotiana t	Ogdzcg mus musculu	Q8i8f3 drosophila	O9D8e5 kluvveromyc	09qvf7 rattus sp.	
036300	Q14182	098905	Q8MJE4	O9TNQ8	Q9QVI3	Q9UCC2	000604	Q16220	OBWIGO	07X566	OBMIO	Q8MJP8	QBMJP7	QBMJP9	Q8MJE3	Q8MJQ1	Q8MJE2	Q9QVK4	Q8R2F6	Q9UJ52	095179	Q26323	P82439	630Z060	Q8I8F3	OSPBES	090VF7	Q9BZ49
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ALIGNMENTS

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0945C1; CTEMBLE 19, Created)
01-DEC-2001 (TEMBLE 19, Last sequence update)
01-DEC-2001 (TEMBLE 1.19, Last sequence update)
01-DEC-2001 (TEMBLE 1.19, Last annotation update)
Dip5 protein (Fragment).
Crypthecodinium cohnii (Dinoflagellate).
Eukaryota; Alveolata; Dinophyceae; Gonyaulacales; Crypthecodiniaceae;
Crypthecodinium.
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SEQUENCE FROM N.A.
MEDLINE=21428164; PubMed=11545436;
Guillebault D., Derelle E., Bhaud Y., Moreau H.;
"Role of nuclear WW domains and proline-rich proteins in dinoflagellate transcription.";
Protist 152:127-138 (2001).
EMBL; AR41750, AAL15908.1;
NON_TER
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01-MAR-2003 (TrEMBLrel. 23, Created)
01-MAR-2003 (TrEMBLrel. 23, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
                                                                                              13 A.A.
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55.6%;
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Best Local Similarity 55.6
Matches 5; Conservative
                                                                                      PRELIMINARY;
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RESULT 1
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                                                                                                                Tran P., Leclerc D., Chan M., Pai A., Hiou-Tim F., Wu Q., Goyette P., Artigas C., Milos R., Rozen R.; "Multiple transcription start sites and alternative splicing in the methylenetrahydrofolate reductase gene result in two enzyme
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Mammalia, Eutheria, Rodentia, Sciurognathi, Muridae, Murinae, Mus.
NCBI_TaxID=10090;
                                Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Rodentia, Sciurognathi, Muridae, Murinae, Mus
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Methylenetetrahydrofolate reductase short isoform (Fragment).
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01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
Methylenetetrahydrofolate reductase short isoform (Fragment).
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Pred. No. 1.5e+03;
0; Mismatches 4; Indels
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Pred. No. 1.4e+03;
0; Mismatches 4; Indels
                                                                                                                                                                                                                        SEQUENCE 14 AA; 1472 MW; 28DD341AC1695CB8 CRC64;
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01-MAY-1999 (TrEMBLrel. 10, Last sequence update)
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MEDLINE=22257759; Pubmed=12370778;
                                                                                                      MEDLINE=22257759; PubMed=12370778;
                                                                                                                                                                            Mamm. Genome 13:483-492(2002).
                                                                                                                                                                                       EMBL; AF404271; AAN40873.1; -. MGD; MGI:106639; Mthfr.
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nilarity 55.6%;
Conservative 0
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55.6%;
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MGD; MGI:106639; Mthfr.
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                      Mus musculus (Mouse)
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Best Local Similarity
Matches 5; Conserv
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Best Local Similarity
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                                                    NCBI_TaxID=10090;
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                                                                                            STRAIN=129/Sv;
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STRAIN=Bassan; TISSUE=Leaf;
MEDLINE=20133023; PubMed=10666448;
Medline=20133023; PubMed=10666448;
del Campo E., Sabater B., Martin M.;
"Transcripts of the ndhH-D operon of barley plastids: possible role of unedited site III in splicing of the ndhA intron.";
Nucleic Acids Res. 28:1092-1098(2000).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SECUENCE FROM N.A. MEDINDE=97019047; PubMed=8865663; Sites J.W. JT., Davis S.K., Guerra T., Iverson J.B., Snell H.L.; Sites J.W. JT., Davis S.K., Guerra T., Iverson J.B., Snell H.L.; "Character congruence and phylogenetic signal in molecular and morphological data sets: a case study in the living Iguanas (Squamata,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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                                                                              Bukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; Pooideae;
Triticeae; Hordeum.
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Lepidosauria, Squamata, Iguania, Iguanidae, Iguaninae, Ctenosaura,
NCBI_TaxID=51212;
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1 1 1 SEQUENCE 14 AA; 1521 MW; 361CF5911A68F8B8 CRC64;
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01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
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Last annotation update)
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Pred. No. 2e+03;
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EMBL; U66228; AAB07474.1; -.
GO; GO:0005739; C:mitochondrion; IEA.
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GO; GO:0009507; C:chloroplast; IEA.
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                      ORF321 protein (Fragment).
Hordeum vulgare (Barley).
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Mol. Biol. Eve
                                                                 Chloroplast
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Q95952;
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                                                                                                                                                                                              Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Lepidosauxia, Squamata; Iguania; Iguanidae; Iguaninae; Sauromalus.
NCBI_TaxID=51220;
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Lepidosauria, Squamata, Iguania, Iguanidae, Phrynosomatinae,
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SEOUENCE 15 AA; 1843 MW; 8327179CE9203901 CRC64;
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01-FEB-1997 (TrEMBLrel. 02, Last sequence update)
01-UNN-2003 (TrEMBLrel. 24, Last annotation update)
                                                                                                                                           Sauromalus varius (San Esteban island chuckwalla).
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EMBL; U66233; AAB07480.1; -
GO; GO:0005739; C:mitochondrion; IEA.
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EMBL, U66224, AAB07478.1, -.
GO; GO:0005739; C:mitochondrion; IEA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Chernajovsky Y., Brown A., Jones T.A.; "Promoter first exon'/intron characterization and chromosomal location of the human light chain (beta) kinesin gene."; DNA Cell Biol. 15:965-974 (1996).
BMBL, X69688, CAA49349.1; -.
PRMBL, S43170, S43170.
PIR, S43170, S43170.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Lepidosauria; Squamata; Iguania; Iguanidae; Iguaninae; Cyclura.
NCBI_TaxID=51215;
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01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
01-FEB-1997 (TrEMBLrel. 02, Created)
1-FEB-1997 (TrEMBLrel. 02, Last sequence update)
01-0CT-2003 (TrEMBLrel. 25, Last annotation update)
NADH dehydrogenses subunit 4 (Fragment).
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                                                                                                                 Cyclura ricordi (Ricord's rock iguana)
                                                                                                                                                                                                                                                                                                                                                              Iguanidae).";
Mol. Biol. Evol. 13:1087-1105(1996).
EMBL; U66237; AAB07473.1; -.
Mitochondrion.
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MEDLINE=97019047; PubMed=8865663;
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BETA-KINESIN.
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                                                                                                                                         Mitochondrion.
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SEQUENCE
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STRAIN=cv. Money Maker; TISSUE=Leaf;
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Mammalia, Eutheria, Rodentia, Sciurognathi, Muridae, Murinae, Rattus.
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NCBI_TaxID=10632;
                                                                                     Anthopleura elegantissima (Sea anemone).
Eukaryota; Metazoa; Cuidaria; Anthozoa; Zoantharia; Actiniaria;
Nynantheae; Actiniidae; Anthopleura.
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MEDLINE=96542418; PubNed=9750786;
Takemura M., Yamauchi K., Yamarodani A.;
"Structural analysis of histamine N-methyltransferase gene.";
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Pred. No. 1e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 23; DB 5; Length 9; Pred. No. 1e+06;
01-MAY-2000 (TrEMBLrel. 13, Last sequence update) 01-MAY-2000 (TrEMBLrel. 13, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      01-WAR-2003 (TrEMBLrel. 23, Created)
01-WAR-2003 (TrEMBLrel. 23, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   9 AA; 1060 MW; 4223B1E042CEB053 CRC64;
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01-OCT-2002 (TrEMBLrel. 22, Created)
01-OCT-2002 (TrEMBLrel. 22, Last sequence update)
01-OCT-2002 (TrEMBLrel. 22, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                anemones.";
Peptides 13:851-857(1992).
CROUINCE 9 AA; 943 MW; 2908176737686777 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Methods Find. Exp. Clin. Pharmacol. 17:1-4 (1995).
MELS, 882579; AANGOT45.1; -.
GOS, GO:0008168; F.methyltransferase activity; IEA.
GO; GO:0016740; F:transferase activity; IEA.
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                                                                                                                                                                                                                                                                                   MEDLINE=93126143; PubMed=1480510;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Transferase; Methyltransferase.
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80.0%;
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                                                           Antho-RPAMIDE=NEUROPEPTIDE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
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A dis-acting regularory element that affects the alternative splicing of a muscle-specific exon in the mouse NCAM gene.";
Biochim. Biophys. Acta 1397:305-315(1998).

EMBL, AB01873; BAA31275.1; -.

NGD, MGI:97281; NCaml.

NON_TER.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bukaryota, Metazoa, Chordata, Craniata, Vertebrata, Buteleostomi,
Mammalia, Butheria, Rodentia, Sciurognathi, Muridae, Murinae, Mus.
                                                                     Bofill-Mas S., Formiga-Cruz M., Clemente-Casares P., Calafell F.,
                                                                                                                     "Potential transmission of human polyomaviruses through the gastrointestinal tract after exposure to virions or viral DNA."; U. Virol. 75:10290-102299(2001).
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01.MAR-2003 (TrEMBLrel. 23, Last sequence update)
01.MAR-2003 (TrEMBLrel. 23, Last annotation update)
WRXY transcription factor IId-1 splice variant 2 (Fragment).
                                                                                                                                                                                                                                                                                                   Score 23; DB 12; Length 10; Pred. No. 3.1e+03; 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  27.4%; Score 23; DB 11; Length 13; 80.0%; Pred. No. 4e+03; tive 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                     SEQUENCE 10 AA; 1183 MW; 4CD6A96731A32763 CRC64;
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Last annotation update)
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                                                                                                                                                                                                                                                                                                                                                      1; Mismatches
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01-NOV-1998 (TIEMBIREL 08, Last sequence
01-CCT-2002 (TIEMBIREL 22, Last annotation
Neural cell adhesion molecule (Fragment).
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                                             MEDLINE=21465052; PubMed=11581397;
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STRAIN-Balb-c; TISSUE=Liver;
MEDLINE=98250618; PubMed=9582442;
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les 4; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                        6 FGSPRPLP 13
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Best Local Similarity
SEQUENCE FROM N.A.
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Hedegaard J., Kristensen J.E., Nakamura Y., Sperling-Petersen H.U.,
Mortensen K.K.;
Cagna G., Boes A., Ulker B., Zhou A., Wanke D., Somssich I.E.; "Comparison of WRKY group II transcription factors from plants."; Submitted (SEP-2002) to the EMBL/GenBank/DDBJ databases. EMBL; ANY1729.1; -. NON_TER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              "Sequence of the infB gene from Escherichia coli strain 10489 and 10499.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Bacteria; Proteobacteria; Gammaproteobacteria; Enterobacteriales;
Enterobacteriaceae; Escherichia.
NCBL_TaxID=562;
                                                                                                                                                  Query Match

26.2%; Score 22; DB 10; Length 8;

Best Local Similarity 80.0%; Pred. No. 1e+06;

Matches 4; Conservative 0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Submitted (FEB-1999) to the EMBL/GenBank/DDBJ databases EMBL; AJ132862; CAC20133.1; -. NON TER 11 11 SEQUENCE 11 AA; 1319 MW; 6B234CFE740879CB CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                    ©9EUZ3;
01-MAR-2001 (TrEMBLrel. 16, Created)
01-MAR-2001 (TrEMBLrel. 16, Last sequence update)
01-MAR-2001 (TrEMBLrel. 16, Last annotation update)
Ribosome binding factor A (Fragment).
                                                                                                    SEQUENCE 8 AA, 917 MW, 59177B4775B87330 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                    11 AA.
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/cgnz_6/ptodata/2/jaa/5B_COMB.pep:*
/cgnz_6/ptodata/2/jaa/6A_COMB.pep:*
/cgnz_6/ptodata/2/jaa/6A_COMB.pep:*
/cgnz_6/ptodata/2/jaa/PcTUS_COMB.pep:*
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GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
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US-08-602-999A-255

US-08-602-999A-410

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Match Length DB
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999 003 224 224 44 44 44 44 44 44 44 44 44 44 4	C-SRC SH3	1; Length 3; Ind
30-047-29 45-632-103 00-12947-355 00-12947-34 52-877-34 76-515A-34 76-515A-34 76-515A-34 76-515A-34 76-515A-34 76-515A-34 76-515A-34 76-515A-34 76-515A-34 76-515A-34 76-515A-34 76-515A-34 76-515A-34 76-615A-34 78-865-85 78-865	MENTS C.	re 37; DB d. No. 21; Mismatches
US-08-230-047-29 US-09-045-632-103 US-08-632-103 US-08-36-315-34-35 US-08-35-877-34 US-08-652-877-34 US-08-652-877-34 US-08-652-877-34 US-08-652-877-34 US-08-652-877-34 US-08-652-877-34 US-08-652-877-34 US-09-500-124-201 US-09-500-124-85 US-09-500-124-85 US-09-500-124-85 US-09-500-124-85 US-09-500-124-85 US-08-650-124-85 US-08-650-124-85 US-08-650-124-85 US-08-650-124-85 US-08-650-124-85 US-08-650-124-85 US-08-650-124-85 US-08-650-124-85 US-08-650-124-85 US-08-650-124-85 US-08-765-783A-83	LUS/08230047 III, George H. Ko, Yuri D. EXPRESSION CLONING OPROTEIN A.4 0.5 coulenc Rorer Inc. Road, 3C43 Road, 3C43 W. disk h. System 7.1 (Patentin) ATA: (Patentin) ATA: 32/84 SATION: Anne 32/534 MBER: Al465-US GWANTION: 54-3817 SATION: 64-3817 SATION:	Score 37 Pred. No 0; Misma
	s/0823006 s/0823006 runi D. C. George Turi D. C. Seorge Tression T	
0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0	Lication US/082300 TON: antobe III, Georg Asshchenko, Yuri II Asshchenko, Yuri II ATION: EXPRESSION THION: EXPRESSION THION: PROTEIN THOSE: 40 TH	44.0% 70.0% ative
CO C	Licat 11001: 110	Similarity 70.
	ULT 1 08-230-047-30 equence 30, Applica equence 30, Applica APPLICANT: Search APPLICANT: Jase, APPLICANT: Jase, APPLICANT: Jase, APPLICANT: Jase, APPLICANT: Jase, APPLICANT: Jase, ADDRESSE: Rho STREET: 500 Arr CITY: Collegev STREET: 500 Arr CITY: Collegev STREET: PA COUNTRY: WAS COUNTRY: WAS COUNTRY: MACINGUM FELLING DATE: 1 CLASSIFICATION NUM FELLING DATE: 1 CLASSIFICATION NUM FELEPRONE: (610) TELEPRAX: (610) TELEPRONE: TYPE: P	atch cal Sim:
2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2	RESULT 1 Sequence 30 Sequence 30 Sequence 30 Sequence 30 Patent No. GENERAL INO. TITLE OF TIT	Query Match Best Local Matches

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TOPOLOGY:
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                                                 COUNTRY:
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Patent No. 6303574
GENERAL INFORMATION
APPLICANT: RAY, BRIAN K.
APPLICANT: SPARKS, ANDREW B.
APPLICANT: THORN, JUDITH M.
APPLICANT: QUILLIAM, LAWRENCE A.
APPLICANT: DER, CHANNING J.
TITLE OF INVENTION: SIC SH3 BINDING PEPTIDES AND METHODS OF TITLE OF INVENTION: ISOLATING AND USING SAME NUMBER OF SEQUENCES: 106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OBLON, SPIVAK, MCCLELLAND, MAIER & NEUSTADT, P.C.
                                                                                                                                                                                                                                                                                                                                                                                                 RIDER, James E. WVENTION: SH3 BINDING PEPTIDES AND METHODS OF VVENTION: ISOLATING AND USING SAME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           41.7%; Score 35; DB 3; Length 13; 62.5%; Pred. No. 36; 0; Indels tive 3; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STATE: New York
COUTRY: U.S.A.
ZIP: 10.36-ZIP: 10.36-ZI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1101-202
                                                                                                                                                                                  APPLICANT: KAY, Brian K.
APPLICANT: THORN, Judith M.
APPLICANT: OULLLIAM, Lawrence A.
APPLICANT: POWLKES, Dana M.
APPLICANT: FOWLKES, Dana M.
APPLICANT: RIDER, James E.
TITLE OF INVENTION: SH3 BINDING PE
TITLE OF INVENTION: ISOLATING AND
NUMBER OF SEQUENCES: 467
                                 Sequence 92, Application US/08602999A Patent No. 6184205
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Misrock, S. Leslie
REGISTRATION NUMBER: 18,872
REFERENCE/DOCKET NUMBER: 1101-:
TELECOMUTUNICATION:
TELEPHONE: (212) 790-9090
TELEFAX: (212) 869-9741/8864
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pennie & Edmonds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEX: 66141 PENNIE INFORMATION FOR SEQ ID NO: 92: SEQUENCE CHARACTERISTICS:
                                                                                                                                                         Andrew B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               13 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSE: Pennie &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              unknown
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2 TPRPLPLI 9
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Best Local Similarity
Matches 5; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-08-602-999A-92
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US-08-602-999A-92
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY:
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Sequence 92, Application US/09500124

Patent No. 6432920

GENERAL INFORMATION:
APPLICANT: SPARKS, Andrew B.
APPLICANT: THORN, Judith M.
APPLICANT: Channing U.
APPLICANT: FOWLKES, Dana M.
APPLICANT: FOWLKES, Dana M.
APPLICANT: FOWLKES, Dana M.
APPLICANT: RIDER, James E.
TITLE OF INVENTION: SB13 BINDING PEPTIDES AND METHODS OF TITLE OF INVENTION: 1SOLATING AND USING SAME
NUMBER OF SEQUENCES: 467
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  41.7%; Score 35; DB 4; Length 13; 62.5%; Pred. No. 36;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Indels
[: 1755 S. Jefferson Davis Highway, Suite 400
Arlington
    Virginia
                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/278,865
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/500,124
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               3; Mismatches
                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
NAME: Villacorta, Gilberto M.
REGISTRATION NUMBER: 34,038
REFERENCE/DOCKET NUMBER: 4980-007-0
TELECOMMUNICATION:
TELEPHONE: (703) 413-3000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ANDRESSEE: Pennie k bumber.
STREET: 1155 Avenue of the Americas
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STATE: New York
COUNTRY: U.S.A.
ZIP: 10036-2711
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPOTER: 1BM PC COMPALIBLE
OPERATING SYSTEM: PC_DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US/09/500,124
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CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/602,999
                                                                                                                                                                                                                                                                                                                                                                                                                           TELEPHONE: (703) 413-300
TELERAX: (703) 413-220
TELEX: 24885 OPAT UR
INFORMATION FOR SEQ ID NO: 9
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 16-FEB-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 13 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 5; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                         FILING DATE:
CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          unknown
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Best Local Similarity
Matches 5; Conserv?
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2 TPRPLPLI 9
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                                                                    U.S.A.
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APPLICANT: Castelhano, Arlindo
APPLICANT: Witter, David
TITLE OF INVENTION: CONFORMATIONALLY CONSTRAINED PEPTIDOMIMETICS AS BETA-TURN
TITLE OF INVENTION: TEMPLATES AND MODULATORS OF SH3 DOMAINS
FILE REFERENCE: 60384-B
CURRENT APPLICATION NUMBER: US/09/086,168B
CURRENT FILING DATE: 1998-05-28
NUMBER OF SEQ ID NOS: 5
SOFTWARE: Patentin version 3.1
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Sequence 29, Application US/08336343A

Patent No. 5677144

GENERAL INFORMATION:

APPLICANT: Ullrich, Axel

APPLICANT: Alves, Frauke

TILLE OF INVENTION: CCK-2, A No. 5677144el Receptor Tyrosine Kinase

NUMBER OF SEQUENCES: 43

CORRESPONDENCE ADDRESS:
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41.7%; Score 35; DB 4; Length 13;
Best Local Similarity 62.5%; Pred. No. 36;
Matches 5; Conservative 3; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     40.5%; Score 34; DB 4; Length 9; 100.0%; Pred. No. 3e+05; tive 0; Mismatches 0; Indels
               NAME: Misrock, S. Leslie
REGISTRATION NUMBER: 18,872
REFERENCE/DOCKET NUMBER: 1101-202
TELEPHONE: (212) 790-9090
TELEPAX: (212) 790-9090
TELERAX: 66141 PENNIE
TELERAX: 66141 PENNIE
TELERAX: 66141 PENNIE
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE: Pennie & Edmonds
STREET: 1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CCATION: (1). (9)
CTHER INFORMATION: Synthetic Peptide
US-09-086-1688-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OTHER INFORMATION: SH3 test peptide FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 3, Application US/09086168B Patent No. 6638941 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: PRT
ORGANISM: Artificial Sequence
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                : 13 amino acids
amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                             , MOLECULE TYPE: peptide US-09-500-124-92
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME/KEY: MISC_FEATURE
                                                                                                                                                                                                                                                           unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                 8 SPRPLPW 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ouery Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2 TPRPLPLI 9
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                                                                                                                                                                                                                                                         TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-09-086-168B-3
                                                                                                                                                                                                            LENGTH:
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LENGTH: 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ద
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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       40.5%; Score 34; DB 1; Length 11; 100.0%; Pred. No. 43; tive 0; Mismatches 0; Indels
                                                                                                                                               SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/336,343A
FILING DATE: 08-NOV-1994
CLASSIPICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSEE: Rhone-Poulenc Rorer Inc.
STREET: 500 Arcola Rd., 3C43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION: 435
PRICR APPLICATION DATA:
APPLICATION NUMBER: PCT/US95/15203
FILING DATE: 22-NOV-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/344,836
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/08/652,877
                                                                                   MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
NAME: CCTUZZI, LBUTZ A.
REGISTRATION NUMBER: 30,742
REPERBUCK/DOCKET NUMBER: 7663-
TELECOMUNICATION INFORMATION:
TELEPHONE: (212) 790-3904
TELEPHONE: (212) 869-9741/8864
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 29, Application US/08652877
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: Macintosh
OPERATING SYSTEM: System 7.5.1
SOFTWARE: Word 6. (Patentin)
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Patent No. 6187548
GENERAL INFORMATION:
APPLICANT: Akerstrom, Goran
APPLICANT: Cimilan, Claes
APPLICANT: Crumley, Gregg R.
APPLICANT: Crumley, Gregg R.
APPLICANT: Murray, Gregg R.
APPLICANT: Murray, Edward M.
APPLICANT: Hjalm, Goran
                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: (212) 869-9741/886
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 29:
SEQUENCE CHARACTERISTICS:
LENGTH: 11 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ZIP: 19426-0107
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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Best Local Similarity 100.
Matches 6; Conservative
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MOLECULE TYPE: peptide
                                       ZIP: 10036-2711
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STREET: 500 Arcola CITY: Collegeville
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               amino acid
New York
U.S.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-08-336-343A-29
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STATE:
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Query Match
Best Local Similarity 100.v
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Best Local Similarity 100.0
Matches 6; Conservative
                                                                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 11 amino acids
                                                                                                                                                                                                                                                                  internal
                                                                                                                                                                                                          TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: NO
                                                                                                                                                                      amino acid
                                                                                                                                                                                                                                                                                                                                                                                                  9 PRPLPV 14
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US-08-476-515A-29
                                                                                                                                                                                      STRANDEDNESS:
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APPLICANT: Juhlin, Claes
APPLICANT: Juhlin, Claes
APPLICANT: Abit, Lars
APPLICANT: Crumley, Gregg R.
APPLICANT: Morse, Clarence C.
APPLICANT: Morray, Edward M.
APPLICANT: Hjalm, Goran
TITLE OF INVENTION: Thereof and DNA Encoding Same
NUMBER OF SEQUENCES: 84
CORRESPONDENCE ADDRESS:
ADDRESSEE: Martin Savitzky
                                                                                                                                                                                                                                                                                                                                                                                                                                                         .
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: Martin Savitzky
STREET: Rhone-Poulenc Rorer Inc., 500 Arcola Rd.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 07-JUN-1995
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/344,836
FILING DATE: 23-NOV-1994
PRIOR APPLICATION NUMBER: W0 PCT/SE94/00483
FILING DATE: 24-MAY-1994
PRIOR APPLICATION DATA:
                                                                                      NAME: SAVIEZRY, MARTIN
REGISTRATION NUMBER: 29,699
REPERENCE/DOCKET NUMBER: 41355E-US
TELEPHONE: 610-454-3816
TELEPHONE: 610-454-3808
INFORMATION FOR SEQ ID NO: 29: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            UMBER: US/08/476,515A
07-JUN-1995
                                     US 08/487,314
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FILING DATE: 24-MAY-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 29, Application US/08476515A Patent No. 6239270 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Compaq PC
OPERATING SYSTEM: Windows 95
SOFTWARE: Word 7.0 (Patentin)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/476,5
FILING DATE: 23-NOV-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/
FILING DATE: 07-UDME-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                               LENGTH: 11 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity 100.
Matches 6; Conservative
                                                                                                                                                                                                                                                                                                                                                             internal
                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE: peptide
HYPOTHETICAL: NO
FRAGMENT TYPE: internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STREET: 3C43,
CITY: Collegeville
STATE: PA
                                                                                                                                                                                                                                                             TYPE: amino acid STRANDEDNESS:
                                                                                                                                                                                                                                                                                                        linear
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                                                                                                                                                                                                                                                                                                      TOPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY:
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REGISTATION NUMBERS: 23,639
REGISTATION NUMBERS: 23,639
REGISTATION NUMBERS: 32,639
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REGISTATION NUMBERS: 32,639
REGISTATION NUMBERS: 32,645300
REGISTATION NUMBERS: 41,645300
REGISTATION 1,1 amino acida
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REGISTATION 1,1 amino acida
REGISTATION REGISTAT
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Query Match
40.5%; Score 34; DB
Best Local Similarity 100.0%; Pred. No. 47;
Matches 6; Conservative 0; Mismatches
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STREET: 1155 Avenue of the Americas
CITY: New York
                                                                                          1101-202
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-08-602-999A-410; Sequence 410, Application US/08602999A; Patent No. 6184205; GENERAL INFORMATION:
                                         NAME: Misrock, S. Leslie
REGISTRATION WUMBER: 1101-2
REFERENCE/DOCKET NUMBER: 1101-2
TELECOMMUNICATION INPORMATION:
TELEPRONE: (212) 869-9741/8864
TELEFAX: (212) 869-9741/8864
TELEFX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 255:
SEQUENCE CHARACTERISTICS:
LENGTH: 12 amino acids
TYPE: amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ATTORNEY AGENT INFORMATION:
NAME: Misrock, S. Leslie
REGISTRATION NUMBER: 18,872
REFERRINCE/DOCKET NUMBER: 1101-:
TELECOMMUNICATION INFORMATION:
TELEFRAX: (212) 790-9090
TELEFAX: (212) 869-9741/8864
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 410:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: 16-FEB-1996
  FILING DATE: 16-FEB-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   15 amino acids
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                                                                                                                                                                                                                                                                                                                , MOLECULE TYPE: peptide US-09-500-124-255
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; MOLECULE TYPE: peptide US-08-602-999A-410
                                                                                                                                                                                                                                                                                                unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                unknown
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                                                                                                                                                                                                                                                                                           TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
APPLICANT: SPARKS, Andrew B.
APPLICANT: KAY, Brian K.
APPLICANT: THORN, Judith M.
APPLICANT: DER, Channing J.
APPLICANT: POWLKS, Dana M.
APPLICANT: RIDER, James B.
ITILE OF INVENTION: SH3 BINING PEPTIDES AND METHODS OF TITLE OF INVENTION: ISOLATING AND USING SAME
NUMBER OF SEQUENCES: 467
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             40.5%; Score 34; DB 3; Length 12; 100.0%; Pred. No. 47; tive 0; Mismatches 0; Indels
                                SITELL I. 122 AVENUE OF THE AMERICAS
CITY: New York
CONTR: New York
CONTR: U.S.A.
ZIP: 10036-2711
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/NS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATE:
APPLICATION NUMBER: US/08/602,999A
FILLING DATE: 16-FEB-1996
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAMM: MASTOCK, S. Leslie
REGISTRATION NUMBER: 1101-202
REFERENCE/DOCKET NUMBER: 1101-202
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
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CITY: New York
STATE: New York
COUNTRY: New York
COUNTRY: U.S.A.
ZIP: 10036-2711
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
"MEDIUM TYPE: Floppy disk
"MEDIUM TYPE: Floppy disk
"MEDIUM TYPE: Floppy disk
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1155 Avenue of the Americas
                       1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/500,124
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 255, Application US/09500124 Patent No. 6432920
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/602,999
                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE: (212) 790-9090
TELEFAX: (212) 869-9741/8864
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 255:
SEQUENCE CHARACTERISTICS:
LENGTH: 12 amino acids
ADDRESSEE: Pennie & Edmonds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 100.u
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; MOLECULE TYPE: peptide US-08-602-999A-255
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US-09-500-124-255
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY:
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Gaps
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APPLICANT: KAY, Brian K.
APPLICANT: THORN, Judith M.
APPLICANT: QUILLIAM, Lawrence A.
APPLICANT: POWLKES, Dana M.
APPLICANT: FOWLKES, Dana M.
APPLICANT: FOWLKES, Dana M.
APPLICANT: STORE, Janes E.
TITLE OF INVENTION: S13 BINDING PEPTIDES AND METHODS OF TITLE OF INVENTION: 1SOLATING AND USING SAME
NUMBER OF SEQUENCES: 467
CORRESPONDENCE ADDRESS:
ADDRESSEE: Pennie
Score 34; DB 4; Length 12; Pred. No. 47;
                                          0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STATE: New York
COUNTRY: U.S.A.
ZIP: 10036-2711
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OFFRATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN PC-DOS/MS-DOS
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/602,999A
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; MOLECULE TYPE: peptide US-08-602-999A-86
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: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Patent No. 6184205
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               10036-2711
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CLASSIFICATION:
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US-08-602-999A-86
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APPLICANT:
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                                                                                                                                                                                                                                                                                                                                SEQ ID NO 76
LENGTH: 11
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                              40.5%; Score 34; DB 3; Length 15; 100.0%; Pred. No. 58; 0; Indels iive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: SPREKS, Andrew B.
APPLICANT: KAY, Brian K.
APPLICANT: THORN, Uddith M.
APPLICANT: OULLLIAM, Lawrence A.
APPLICANT: DER, Channing J.
APPLICANT: FOWLKES, Dana M.
APPLICANT: FOWLKES, Dana M.
APPLICANT: APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             40.5%; Score 34; DB 4; Length 15; 100.0%; Pred. No. 58; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/500,124
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET: 1155 Avenue of the Americas CITY: New York STATE: New York COUNTRY: U.S.A. ZIP: 10036-2711 COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1101-202
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/602,999
FILING DATE: 16-FEB-1996
ATTORNEY/AGENT INFORMATION:
NAME: Misrock, S. Leslie
REGISTRATION NUMBER: 18,972
REFERENCE/DOCKET NUMBER: 1101-203
TELECOMMUNICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC_compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; Sequence 76, Application US/09417608A; Patent No. 6686164; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                          Sequence 410, Application US/09500124 Patent No. 6432920
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (212) 790-9090
(212) 869-9741/8864
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 410:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pennie & Edmonds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity 100.0
-hea 6; Conservative
Query Match
Best Local Similarity 100..
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
ADDRESSEE: Pennie &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid
TOPOLOGY: unknown
                                                                                                                                                                 9 PRPLPV 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                9 PRPLPV 14
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                                                                                                                                                                                                                                                                                                                                                               US-09-500-124-410
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US-09-417-608A-76
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: RIDER, James E. TITLE OF INVENTION: SH3 BINDING PEPTIDES AND METHODS OF TITLE OF INVENTION: ISOLATING AND USING SAME NUMBER OF SEQUENCES: 467
                                                                                                                                                                                                                                                                                                                                                                                                         3; Indels
APPLICANT: Roggen, Erwin
APPLICANT: Ernst, Steffen
TITLE OF INVENTION: Low Allergenic Protein Variants
FILE REPERENCE: 5676.200-US
CURRENT APPLICATION NUMBER: US/09/417,608A
CURRENT FILING DATE: 1999-10-13
NUMBER OF SEQ ID NOS: 101
SOFTWARE: PatentIn version 3.2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OPERATION SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/602,999A FILING DATE: 16-FEB-1996
                                                                                                                                                                                                                                                                                                                                                                                                         1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE: Pennie & Edmonds
STREET: 1155 Avenue of the Americas
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1101-202
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SPARKS, Andrew B.
KAY, Brian K.
THORN, Judith M.
QUILLIAM, Lawrence A.
DER, Channing J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 86, Application US/08602999A Patent No. 6184205
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 790-9090
TELEFAX: (212) 869-9741/8864
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           18,872
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INFORMATION FOR SEQ ID NO: 86:
SEQUENCE CHARACTERISTICS:
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Best Local Similarity 60.0%;
Matches 6; Conservative
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NAME: Misrock, S. Leslie
REGISTRATION NUMBER: 18, R
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                            CTHER INFORMATION: Synthetic US-09-417-608A-76
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 13 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 CRAHSFGSPR 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      unknown
                                                                                                                                                                                                                                               ORGANISM: Artificial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 amino acid
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0
                                               0; Gaps
                                           0; Indels
Query Match 39.3%; Score 33; DB 3; Length 13; Best Local Similarity 62.5%; Pred. No. 71; Matches 5; Conservative 3; Mismatches 0; Indels
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8 SPRPLPVV 15 :|||||: 2 TPRPLPML 9

Pp.

Search completed: April 2, 2004, 10:12:44 Job time : 14 secs

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GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
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OM protein - protein search, using sw model

April 2, 2004, 10:11:14; Search time 32.3333 Seconds (without alignments) 121.720 Million cell updates/sec Run on:

US-10-009-049-1 Title: Perfect score:

1 CRAHSFGSPRPLPVV 15 Sequence:

BLOSUM62 Scoring table: 1070241 segs, 262374223 residues Searched:

Gapop 10.0 , Gapext 0.5

Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 15

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database :

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

	Description	Sequence 92, Appl	Seguence 92,	Sequence 17,	0,		410 Sequence 410, App	(C)		S			8 Sequence 8, Appli		S	-
	ID	US-09-938-315-92	US-10-161-791-	US-09-945-917-17	US-09-845-917A	US-10-161-791-	US-10-161-791-410	US-09-938-315-86	US-10-161-791-	US-09-938-315-6	US-10-161-791-	US-10-161-791-	US-10-059-261-	US-10-059-261-116	US-09-938-315-67	73-107-131-01-SI
	DB	6	14	10	10	14	14	σ	14	σ	14	14	14	14	σ	14
	Query Match Length DB	13	13	11	11	12	15	13	13	14	14	15	15	15	14	1.4
æ	Query Match	41.7	41.7	40.5	40.5	40.5	40.5	39.3	39.3	39.3	39.3	38.7	38.1	38.1	36.9	36 9
	Score	35	35	34	34	34	34	33	33	33	33	32.5	32	32	31	71
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TELERAX: (703) 413-2220 TELEX: 248855 OPAT UR

Sequence 155, App Sequence 114, App Sequence 4631, Ap Sequence 30, Appl Sequence 281, Appl	85, i 85, 37, 60,	256 264 301 33,	0 138 0 138	or	Sequence 14, Appl Sequence 14, Appl Sequence 59, Appl Sequence 223, App Sequence 26, Appl Sequence 26, Appl Sequence 27, Appl Sequence 28, Appl
4 US-10-161-7 US-09-815-83 5 US-10-149-1 4 US-10-286-4 4 US-10-161-7	.938-315-85)-161-791-8)-966-459A-	4 US-10-001-469 4 US-10-001-469 4 US-10-161-791 US-09-730-857-	2 US-10-601-837 0 US-09-572-404 0 US-09-572-404 0 US-09-572-404	4 US-10-044-8 0 US-09-291-4 0 US-09-291-4 4 US-10-158-5 4 US-10-158-5	14 US-10-254-440A-14 14 US-10-155-833B-14 14 US-10-193-477-59 14 US-10-161-791-306 14 US-10-008-554A-26 14 US-10-008-524A-28 14 US-10-008-524A-28
1221	11 11 13 14 14	155 155 9	6016	22222	117 117 118 118 118 118
36.9 35.7 35.7 35.7	35.7 35.7 35.7	35.7 35.7 34.5	ይ ይ ይ ይ 4 4 4 4 4 . የ የ የ የ የ የ	ლ თ თ თ თ ს 4 4 4 4 4 4 4 ი ი ი ი ი ი ი	,
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ALIGNMENTS

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APPLICANT: KAY, BRIAN K.
SPARKS, ANDREW B.
THORN, JUDITH M.
QUILLIAM, LAWRENCE A.
DER, CHANNING J.
TITLE OF INVENTION: SIC SH3 BINDING PEPTIDES AND METHODS OF INUMBER OF SEQUENCES: 106
                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
ADDRESSEE: OBLON, SPIVAK, MCCLELLAND, MAIER & NEUSTADT,
                                                                                                                                                                                                                                                                                                                                                                                                    STREET: 1755 S. Jefferson Davis Highway, Suite 400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/938,315
PILING DATE: 23-Aug-2001
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
NAME: Villacorta, Gilberto M.
REGISTRATION NUMBER: 34,038
REFERENCE/DOCKET NUMBER: 4980-007-0
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703) 413-3000
                                              ; Sequence 92, Application US/09938315; Patent No. US20020091085A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ZIP: 22202
COMPUTER READABLE FORM:
MEDIUM TYPE: Flopp)
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STATE: Virginia
COUNTRY: U.S.A.
RESULT 1
US-09-938-315-92
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Pred. No. 1.3e+02;
3; Mismatches 0; Indels
                                                                                                                                                                                   Query Match 41.7%; Score 35; DB 9; Length 13; Best Local Similarity 62.5%; Pred. No. 1.3e+02; Matches 5; Conservative 3; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: STARKS, Andrew B.
APPLICANT: KAY, Brian K.
APPLICANT: KAY, Brian K.
APPLICANT: OUILLIAM, Lawrence A.
APPLICANT: OUILLIAM, Lawrence A.
APPLICANT: DER, Channing J.
APPLICANT: FUBER, James E.
TITLE OF INVENTION: SH3 BINDING PEPTIDES AND METHODS OF
TITLE OF INVENTION: ISOLATING AND USING SAME
NUMBER OF SEQUENCES: 467
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
                                                              TOPOLOGY: unknown

HOLECULE TYPE: peptide

SEQUENCE DESCRIPTION: SEQ ID NO: 92:
US-09-938-315-92
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           E: Pennie & Edmonds
1155 Avenue of the Americas
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APPLICATION NUMBER: US/08/602,999
FILING DATE: 16-FEB-1996
ATTORNEY/AGENT INFORMATION:
NAME: Misrock, S. Leslie
REGISTRATION NUMBER: 18,872
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REGISTRATION NUMBER: 18,872
REFERENCE/DOCKET NUMBER: 1101-202
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 790-9090
TELEFAX: (212) 869-9741/8864
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/161,791
                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 92, Application US/10161791
Publication No. US20030186863A1
GENERAL INFORMATION:
                                          LENGTH: 13 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEX: 66141 PENNIE INFORMATION FOR SEQ ID NO: 92:
INFORMATION FOR SEQ ID NO: 92: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         h
Similarity 62.5%;
5; Conservative 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE: peptide
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ADDRESSEE: Pennie &
                                                                                                                                                                                                                                                                            8 SPRPLPWV 15
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2 TPRPLPLI 9
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RY: U.S.A.
10036-2711
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Best Local Similarity
Matches 5; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                  US-10-161-791-92
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-10-161-791-92
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STATE:
                                                                                                                                                                                                                                                                                                                                                                                  RESULT 2
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                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: JOSECT JULIANT JOSECT JULIANT JOSECT JULIANT JOSECT JULIANT JOSECT JULIANT 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Bogaert, Thierry
APPLICANT: Vandekerckhove, Joel
TITLE OF INVENTION: PROCESSES FOR THE IDENTIFICATION OF COMPOUNDS WHICH
TITLE OF INVENTION: CONTROL CELL BEHAVIOUR, THE COMPOUNDS IDENTIFIED AND
TITLE OF INVENTION: PHARMACEUTICAL COMPOSITIONS CONTIANING THEM AND THEIR
FILLE OF INVENTION: USE IN THE CONTROL OF CELL BEHAVIOUR
FILLS REFERENCE: P/14-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gape
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40.5%; Score 34; DB 10; Length 11;
Best Local Similarity 100.0%; Pred. No. 1.6e+02;
Matches 6; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           40.5%; Score 34; DB 10; I 100.0%; Pred. No. 1.6e+02; tive 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 17, Application US/09845917A Publication No. US20030167538A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; Sequence 255, Application US/10161791; Publication No. US20030186863A1 GENERAL INFORMATION: APPLICANT: SPARKS, Andrew B.
                                                                                                                                                                                                                                                                Sequence 17, Application US/09945917, Publication No. US2030042381A1, GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Caenorhabditis elegans
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: Patentin Ver. 2.1
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2 TPRPLPLI 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2 PRPLPV 7
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                                                                                                                                                                              RESULT 3
US-09-945-917-17
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LENGTH: 11
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RESULT 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
APPLICANT: SPARKS, Andrew B.
APPLICANT: KAY, Brian K.
APPLICANT: THORN, Judith M.
APPLICANT: THORN, Judith M.
APPLICANT: DER, Channing J.
APPLICANT: PUBER, James E.
TITLE OF INVENTION: SH3 BINDING PEPTIDES AND METHODS OF
TITLE OF INVENTION: ISOLATING AND USING SAME
NUMBER OF SEQUENCES: 467
                                                                          APPLICANT: RIDER, James E.
IIILE OF INVENTION: SH3 BINDING PEPTIDES AND METHODS OF
IIILE OF INVENTION: ISOLATING AND USING SAME
                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/161,791
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEE: Pennie & Edmonds
: 1155 Avenue of the Americas
New York
                                                                                                                                    NUMBER OF SEQUENCES: 467
CORRESPONDENCE ADDRESS:
ADDRESSEE: Pennie & Edmonds
STREET: 1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/602,999
FILING DATE: 16-FEB-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1101-202
                                                                                                                                                                                                                                                           COUNTRY: U.S.A.
ZIP: 10036-2711
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMpatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: MISTOCK, S. LEBLIE
REGISTRATION NUMBER: 18,872
REFERENCE/DOCKET NUMBER: 1101-:
TELECOMMUNICATION INFORMATION:
TELEFRONE: (212) 900-9090
TELEFAX: (212) 869-9741/8864
              OUILLIAM, Lawrence A. DER, Channing J. FOWLKES, Dana M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 255:
SEQUENCE CHARACTERISTICS:
THORN, Judith M.
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Matches 6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY: unknown
MOLECULE TYPE: peptide
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                                                                                                                                                                                                                       New York
New York
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                 APPLICANT:
APPLICANT:
                                                       APPLICANT:
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ADDRESSEE: OBLON, SPIVAK, MCCLELLAND, MAIER & NEUSTADT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: KAY, BRIAN K.
SPARKS, ANDREW B.
SPARKS, AUDITH M.
QUILLIAM, LAWRENCE A.
DER, CHANNING J.
TITLE OF INVENTION: Src SH3 BINDING PEPTIDES AND METHODS OF
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 40.5%; Score 34; DB 14; Length 15; 100.0%; Pred. No. 2.1e+02; tive 0; Mismatches 0; Indels
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/938,315
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRANT APPLICATION DATA:
APPLICATION NUMBER: US/10/161,791
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: 23-Aug-2001
CLASSIFFCATION: «Unknown»
ATTORNEY/AGENT INFORMATION:
NAME: Villacorta, Gilberto M.
REGISTRATION NUMBER: 34,038
REFERENCE/DOCKET NUMBER: 4980-007-0
TELECOMMUNICATION INFORMATION:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                             FILING DATE:
CLASSIFICATION:
PRIOR PAPELICATION:
PRIOR APPLICATION NUMBER: US/08/602,999
FILING DATE: 16-FEB-196
ATTORNEY/AGENT INFORMATION:
NAME: Misrock, S. Lesing, REGISTRATION NUMBER: 18,872
REFERRNCE/DOCKET NUMBER: 1101-202
TELECHOME: (212) 790-9090
TELEFRAX: (212) 869-9741/8864
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 86, Application US/09938315
Patent No. US20020091085A1
GENERAL INFORMATION:
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TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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STATE: Virginia
COUNTRY: U.S.A.
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MOLECULE TYPE: peptide
US-10-161-791-410
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Best Local Similarity
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                                                                                                                                                                                                                             Query Match
Best Local Similarity 62.5%; Pred. No. 2.6e+02;
Matches 5; Conservative 3; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: KAY, Brian K.
APPLICANT: THORN, Judith M.
APPLICANT: QUILLIAM, Lawrence A.
APPLICANT: DER, Channing J.
APPLICANT: FUNKES, Dana M.
APPLICANT: RIDER, James E.
TITLE OF INVENTION: ISOLATING PRPTIDES AND METHODS OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/161,791
                                                                                                                                    TOPOLOGY: unknown
MOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: SEQ ID NO: 86:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3: Pennie & Edmonds
1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Misrock, S. Leslie
REGISTRATION NUMBER: 18,872
REFERENCE/DOCKET NUMBER: 1101-202
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 790-9090
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/602,999
FILING DATE: 16-FEB-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
  (703) 413-3000
                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 86, Application US/10161791
Publication No. US20030186863A1
                TELEFAX: (703) 413-2220
TELEX: 24885 OPAT UR
INFORMATION FOR SEQ ID NO: 86:
SEQUENCE CHARACTERISTICS:
LENGTH: 13 amino acids
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62.5%;
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APPLICANT: SPARKS, Andrew B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Floppy disk
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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                                                                                                                 TYPE: amino acid
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CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      unknown
                                                                                                                                                                                                                                                                                                              8 SPRPLPVV 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM PC
OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    amino acid
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2 TPRPLPML 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
    TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New York
: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      10036-2711
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: New York STATE: New Yor!
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 8
US-10-161-791-86
                                                                                                                                                                                               US-09-938-315-86
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Gaps
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ADDRESSEE: OBLON, SPIVAK, MCCLELLAND, MAIER & NEUSTADT,
                                                                                                                                                                                                                                                     APPLICANT: KAY, BRIAN K.

SPARKS, ANDREW B.
THORN, JUDITH M.
QUILLIAM, LAWRENCE A.
DER, CHANNING J.
TITLE OF INVENTION: SIC SH3 BINDING PEPTIDES AND METHODS OF ISOLATING AND USING SAME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ..
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           P.C.
STREET: 1755 S. Jefferson Davis Highway, Suite 400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 33; DB 9; Length 14;
Pred. No. 2.8e+02;
3; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
  ..
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REFERENCE/DOCKET NUMBER: 4980-007-0 TELECOMMUNICATION INFORMATION: TELEPHONE: (703) 413-3000
  3; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/938,315
FILING DATE: 23-Aug-2001
CLASSIFICATION: «UDKNOWN»
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: unknown

MOLECULE TYPE: peptide

SEQUENCE DESCRIPTION: SEQ ID NO: 62:
US-09-938-315-62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Villacorta, Gilberto M. REGISTRATION NUMBER: 34,038
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 62, Application US/10161791;
Publication No. US20030186863A1
GENERAL INFORMATION:
APPLICANT: SPARKS, Andrew B.
APPLICANT: THORN, Judith M.
APPLICANT: QUILLIAM, Lawrence A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             THORN, Judith M.
QUILLIAM, Lawrence A.
DER, Channing J.
                                                                                                                                                                                           Sequence 62, Application US/09938315
Patent No. US20020091085A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: (703) 413-2220
TELEX: 24885 OPAT UR
INFORMATION FOR SEQ ID NO: 62:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 14 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              39.3%;
                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES: 106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CITY: Arlington
STATE: Virginia
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 39.3
Best Local Similarity 62.5
Matches 5; Conservative
    5; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   8 SPRPLPVV 15
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                                            8 SPRPLPW 15
                                                                       2 TPRPLPML 9
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RESULT 13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
APPLICANT: SPARKS, Andrew B.
APPLICANT: KAY, Brian K.
APPLICANT: THORN, Judith M.
APPLICANT: THORN, Judith M.
APPLICANT: POWLKES, Dana M.
APPLICANT: FOWLKES, Dana M.
APPLICANT: FOWLKES, Dana M.
APPLICANT: RIBER, James B.
TITLE OF INVENTION: SH3 BINDING PEPTIDES AND METHODS OF
APPLICANT: FOWLKES, Dana M.
APPLICANT: RIDER, James E.
TITLE OF INVENTION: SH3 BINDING PEFTIDES AND METHODS OF
TITLE OF INVENTION: ISOLATING AND USING SAME
                                                                                                                                                                            STATE: New York
COUNTRY: U.S.A.
ZIP: 10036-2711
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/WS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/161,791
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Misrock, S. Lealle
REGISTRATION NUMBER: 18,872
REFERENCE/DOCKET NUMBER: 1101-202
RELEPRONCATION INFORMATION:
TELEPHONE: (212) 790-9090
TELERAX: (212) 790-9090
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 62:
SEQUENCE CHARACTERISTICS:
LENGTH: 14 amino acids
TYPE: amino acids
                                                                         NUMBER OF SEQUENCES: 467
CORRESPONDENCE ADDRESS:
ADDRESSEE: Pennie & Edmonds
STREET: 1155 Avenue of the Americas
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES: 467
CORRESPONDENCE ADDRESS:
ADDRESSEE: Pennie & Edmonds
STREET: 1155 Avenue of the Americas
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/602,999
FILING DATE: 16-FEB-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; Sequence 378, Application US/10161791; Publication No. US20030186863A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    39.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 10036-2711
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 62.5%,
-hos 5; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; MOLECULE TYPE: peptide US-10-161-791-62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  8 SPRPLPVV 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3 TPRPLPML 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New York
: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-10-161-791-378
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY:
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FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: vector pACgp67-SCFv461
OTHER INFORMATION: peptide sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: EDELMAN, LENA
APPLICANT: BELLMAN, LENA
APPLICANT: BACOTOT.
TITLE OF INVENTION: CHIMERIC MOLECULES CONTAINING A MODULE ABLE TO TARGET
TITLE OF INVENTION: CHIMERIC MOLECULES CONTAINING A MODULE ABLE TO TARGET
TITLE OF INVENTION: PUNCTION OF THE PERMEABILITY TRANSITION PORE COMPLEX
TITLE OF INVENTION: (PTPC)
FILE REFERBNCE: 03495.0216
CURRENT APPLICATION NUMBER: US/10/059,261
CURRENT APPLICATION NUMBER: 00/265,594
PRIOR APPLICATION NUMBER: 60/265,594
PRIOR PILING DATE: 2001-02-02
SOFTWARE: PATENTIN VOS: 325
SOFTWARE: PATENTIN VOS: 325
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               .
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              38.1%; Score 32; DB 14; Length 15; 50.0%; Pred. No. 4.3e+02; tive 2; Mismatches 3; Indels
                                         SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/161,791
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               3;
                                                                                                                               CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/602,999
FILING DATE: 16-FEB-1996
ATTORNEY/AGENT INFORMATION:
NAME: Misrock, S. Leslie
REGISTRATION NUMBER: 18,872
                                                                                                                                                                                                                                                                                                           1101-202
COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 8, Application US/10059261; Publication No. US20030077826A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                TELEPHONE: (212) 790-9090
TELEFAX: (212) 869-9741/8864
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 378:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER: 11(
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 66.7
Matches 8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity 50.0
Matches 5; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 CRAHSFGSPR 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               amino acid
GY: unknown
                                                                                                                   FILING DATE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-10-161-791-378
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-10-059-261-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-10-059-261-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQ ID NO 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
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US-09-938-315-67
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US-10-059-261-116

Sequence 116, Application US/10059261

Sublication No. US20030077826A1

Sublication No. US20030077826A1

SUBLICATION OF US PRIMARY IN THANK INTERIATION TO THE SUBLICANT: EDELMAN, LENA

APPLICANT: EDELMAN, LENA

APPLICANT: BIRAND, JEAN-PAUL

TITLE OF INVENTION: SPECIFIC CELLS AND A MODULE ABLE TO TARGET

TITLE OF INVENTION: FUNCTION OF THE PERMEABILITY TRANSITION PORE COMPLEX

TITLE OF INVENTION: (PPPC)

TITLE OF INVENTION: (PPPC)

TITLE OF INVENTION: (PPPC)

TITLE OF INVENTION: (PPC)

TITLE OF INVENTION: (PPC)

FILE REFERENCE: 03495.0216

CURRENT APPLICATION NUMBER: US/10/059,261

CURRENT FILING DATE: 2001-02-02

NUMBER OF SEQ ID NOS: 325

SOFTWARE: PATENTING USE: 2.1

SEQ ID NO 116

LENGTH: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OTHER INFORMATION: Description of Artificial Sequence: vector pACgp67-SCFv350 OTHER INFORMATION: peptide sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RAL INFURMALIUM.
APPLICANT: KAY, BRIAN K.
SPARKS, ANDREW B.
THORN, UDITH M.
QUILLIAM, LAWRENCE A.
DER, CHANNING J.
TITLE OF INVENTION: Src SH3 BINDING PEPTIDES AND METHODS OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
ADDRESSEE: OBLON, SPIVAK, MCCLELLAND, MAIER & NEUSTADT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      .
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  38.1%; Score 32; DB 14; Length 15; 50.0%; Pred. No. 4.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/938,315
FILING DATE: 23-Aug-2001
CLASSIFICATION: <unknown>
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Villacorta, Gilberto M.
REGISTRATION NUMBER: 34,038
REFERENCE, DOCKET NUMBER: 4980-007-0
TELECHOWICATION INFORMATION:
TELEPHONE: (703) 413-3000
TELEFAX: (703) 413-220
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ZIP: 22202
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 67, Application US/09938315; Patent No. US20020091085A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: PRT ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 50.0%
Local Si Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CITY: Arlington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STATE: Virginia COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 CRAHSFGSPR 10
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                                                                                                                                                                                                                                Length 14;
                                                                                                                                                                                                                             Query Match 36.9%; Score 31; DB 9; Length 14; Best Local Similarity 83.3%; Pred. No. 5.6e+02; Matches 5; Conservative 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: KAY, Brian K.
APPLICANT: THORN, Judith M.
APPLICANT: QUILLIAM, Lawrence A.
APPLICANT: QUILLIAM, Lawrence A.
APPLICANT: FOWLKES, Dana M.
APPLICANT: RIDER, James B.
TITLE OF INVENTION: SH3 BINDING PEPTIDES AND METHODS OF TITLE OF INVENTION: 1SOLATING AND USING SAME
NUMBER OF SEQUENCES: 467
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ZIP: 10036-2711
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/161,791
                                                                                                                TOPOLOGY: unknown
MOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: SEQ ID NO: 67:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        E: Pennie & Edmonds
1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/602,999
FILING DATE: 16-FEB-1996
ATTORNEY/AGENT INFORMATION:
NAME: MISTOCK, S. Leslie
REGISTRATION NUMBER: 18,872
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1101-202
                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-10-161-791-67; Sequence 67, Application US/10161791; Sequence 67, Application US/10161791; Publication No. US20030186863A1; CENERAL INFORMATION: APPLICANT: SPARKS, Andrew B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEPHONE: (212) /20 CT | TELEPHONE: (212) 869-9741/8864
                                                                 LENGTH: 14 amino acids TYPE: amino acid
TBLEX: 248855 OPAT UR INFORMATION FOR SEQ ID NO: 67: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REFERENCE/DOCKET NUMBER: 110
TELECOMMUNICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: (212) 869-9741/886
TELEX: 66141 PENNIE
INPORMATION FOR SEQ ID NO: 67:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 14 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STREET: 1155...
CITY: New York
STATE: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   unknown
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY:
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8 SPRPLP 13 :||||| 3 TPRPLP 8

Sy do

Search completed: April 2, 2004, 10:23:35 Job time: 33.3333 secs

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GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
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OM protein - protein search, using sw model

April Run on:

2, 2004, 09:56:18; Search time 46 Seconds (without alignments) 92.135 Million cell updates/sec

US-10-009-049-2 Perfect score: Title:

1 SRAHSFGSPRPLPVV 15 Sequence: BLOSUM62 Gapop 10.0 , Gapext 0.5 Scoring table:

1586107 segs, 282547505 residues Searched:

Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 15

100% % Post-processing: Minimum Match Maximum Match

Listing first 45 summaries

geneseqp2001s:* geneseqp2002s:* geneseqp2003as:* geneseqp2003bs:* A Geneseq 29Jan04:* geneseqp1980s:* geneseqp1990s:* geneseqp2004s:* geneseqp2000s:* Database

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

STIMMARTES

,		ф			SUMMARIES	
Result No.	Score	Query	Query Match Length	DB	ID	Description
Н	79	100.0	15	4	AAB59250	Aab59250 Peptide #
~	75	94.9	15	4	AAB59249	
m	99	83.5	15	4	AAB59251	
4	38	48.1	15	9	ABR44350	
'n	37	46.8	15	ď	AAW05069	
φ	36	45.6	15	ιΩ	ABG72660	
7	35	44.3	13	~	AAW11111	_
œ	34	43.0	σ	7	AAW94619	Aaw94619 Src homol
თ	34	43.0	15	7	AAW39011	Aaw39011 Peptide r
10	33	41.8	13	7	AAW11105	Ŋ
11	33	41.8	14	N	AAW11098	Src
12	33	41.8	14	~	AAW25512	
13	33	41.8	14	4	AAM53295	Aam53295 Human non
14	32	ö	10	4	AAM06985	Aam06985 CYP1B1 HL
15	32		10	4	AAM08985	HLA-A *
16	32	•	10	4	AAM12950	Aam12950 HLA-A26 d
11	32	6	10	4	AAM08833	Aam08833 HLA-A1 de
8	32	ö	10	4	AAM09489	Aam09489 HLA-B *07
13	32		11	N	AAU75613	Aau75613 Human non
20	32	•	12	m	AAB21272	Aab21272 ADAM 10/K
21	32	•	14	Н	AAP60604	Aap60604 Seguence
22	31	ę.	10	~	AAR93295	
23	31	ę.	10	7	AAR93312	Aar93312 Pi3K prot
24	31	39.5	14	N	AAW11099	σ
25	31		14	7	AAW25517	7 SH3

Aam98385 Human pep	Aam98384 Human pep	Aaw05068 Proline-r	Abb56489 Human sin	Aam13030 HLA-A *02	Aam11185 HLA-A *02	Aam10775 HLA-B8 oc	Aam09513 HLA-B8 oc	Aam08460 HLA-B *27	Aam12240 HLA-B8 no	Aam11714 HLA-A26 n	Aam12099 HLA-A *02	Aam08694 HLA-B *27	Aam11998 HLA-B *15	Aam11458 HLA-A26 n	Aam12119 HLA-A26 n	Aam08347 HLA-Al no	Aam11149 HLA-B8 no		Aam07040 HLA-B *07
AAM98385	AAM98384	AAW05068	ABB56489	AAM13030	AAM11185	AAM10775	AAM09513	AAM08460	AAM12240	AAM11714	AAM12099	AAM08694	AAM11998	AAM11458	AAM12119	AAM08347	AAM11149	AAM07735	AAM07040
4	4	N	4	4	4	4	4	4	4	4	4	4	4	4	4	4	4	4	4
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39.2	39.2	39.5	38.0	38.0	38.0	38.0	38.0	38.0	38.0	38.0	38.0	38.0	38.0	38.0	38.0			38.0	38.0
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26	27	28	29	30	31	32	33	34	35	36	37	38	38	40	41	42	43	44	45

ALIGNMENTS

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New peptides capable of modulating a fibrin fragment E activity, useful for inhibiting uncontrolled cell proliferation, e.g. for treating cancer, vascular restenosis, atherogenesis or psoriasis, and for wound healing.
                                                                       Fibrin fragment E; cell proliferation; cancer; vascular restenosis; arthritis; diabetes; renal disease; psoriasis.
                                                      Peptide #2 recognised by gene VIII library clone 45.
                                                                                                                                                                                                                     ĕ
AAB59250 standard; peptide; 15 AA.
                                                                                                                                                                                                                     Stirk
                                                                                                                                                            07-JUN-2000; 2000WO-GB002197.
                                    (first entry)
                                                                                                                                                                                                                    Thompson WD,
                                                                                                                                                                                                (UYAB-) UNIV ABERDEEN.
                                                                                                                                                                                                                                      WPI; 2001-080581/09.
                                                                                                                       WO200075175-A1.
                                                                                                                                                                             07-JUN-1999;
                                                                                                                                           14-DEC-2000.
                                    27-MAR-2001
                                                                                                                                                                                                                   Melvin WT,
                                                                                                      Synthetic.
                  AAB59250;
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Sequence 15 AA;

Query Match

DB 4; Length 15; 100.0%; Score 79;

The present invention relates to a peptide or its fragment, which is capable of modulating a fibrin fragment E activity. The peptides and modulators are useful for inhibiting stimulation of cell proliferation induced by a fibrin degradation product. The antibody or binding fragment is also useful for identifying the active site of the fibrin fragment receptor. The peptides are particularly useful in treating cancer, vascular restenosis, rhewmatoid archirits, diabetes, renal diseases or psoriasis. The antibodies are also useful for modulating the activity of

Claim 1; Page 49; 59pp; English.

RESULT 2 AAB59249

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New peptides capable of modulating a fibrin fragment E activity, useful for inhibiting uncontrolled cell proliferation, e.g. for treating cancer, vascular restenosis, atherogenesis or psoriasis, and for wound healing.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention relates to a peptide or its fragment, which is capable of modulating a fibrin fragment B activity. The peptides and modulators are useful for inhibiting stimulation of cell proliferation induced by a fibrin degradation product. The antibody or binding fragmen is also useful for identifying the active site of the fibrin fragment B receptor. The peptides are particularly useful in treating cancer, vascular restenosis, rheumatoid arthritis, diabetes, renal diseases or psoriasis. The antibodies are also useful for modulating the activity of
                 Fibrin fragment B; cell proliferation; cancer; vascular restenosis; arthritis; diabetes; renal disease; psoriasis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            N-terminal sequence for guanosine triphosphate activator 10.01.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Guanosine triphosphatase activator 10.01; squamobasal cell;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 66, DB 4; Length 15;
Pred. No. 0.0013;
0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      carcinoma of skin; osteosarcoma; leukemia; teratoma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (SHAN-) SHANGHAI BIOWINDOW GENE DEV INC.
                                                                                                                                                                                                                                                                                                         Ð,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABR44350 standard; peptide; 15 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 1; Page 49; 59pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             83.5%;
92.9%;
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                                                                                                                                                                                             07-JUN-2000; 2000WO-GB002197.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          10-APR-2001; 2001CN-00105912.
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es 13; Conservative
                                                                                                                                                                                                                                                                                                           Thompson WD,
                                                                                                                                                                                                                                                                   (UYAB-) UNIV ABERDEEN
                                                                                                                                                                                                                                                                                                                                               WPI; 2001-080581/09.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 15 AA;
                                                                                                               WO200075175-A1.
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                                                                                                                                                                                                                                 07-JUN-1999;
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                                                                                                                                                                                                                                                                                                         Melvin WT,
                                                                           Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   for inhibiting uncontrolled cell proliferation, e.g. for treating cancer, vascular restenosis, atherogenesis or psoriasis, and for wound healing.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New peptides capable of modulating a fibrin fragment E activity, useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                       Gaps
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                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Peptide #3 recognised by gene VIII library clone 45.
                                                                                                                                                                                                                                                                                                       Peptide recognised by gene VIII library clone 45.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                4.9e-05;
   Pred. No. 1.1e-05; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 75; DB 4;
Pred. No. 4.9e-05
                                                                                                                                                                                                                                                                                                                                                                 arthritis; diabetes; renal disease; psoriasis.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Stirk CM;
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                                                                                                                                                                                             AAB59249 standard; peptide; 15 AA.
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100.0%; Pre
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                       .
100.08;
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                                                           1 SRAHSFGSPRPLPVV 15
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                         15; Conservative
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 Best Local Similarity
Matches 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 15 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                           WO200075175-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           07-JUN-1999;
                                                                                                                                                                                                                                                                       27-MAR-2001
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                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic.
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Best Local (
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Matches

δ g RESULT 3

AAB5925

.; 0

Gaps

.; 0

WPI; 2003-222552/22

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A fusion phage library for expressing random 15mer peptides was prepared using the filamentous bacteriophage fuse 5B. The display library was screened for peptides which bind to the Src SH3 domain. A sequence comparison of the positive peptides revealed a highly reiterated, proline rich, 7 amino acid consensus motif (i.e. the sequence RPLPXXP). This motif demonstrated a strong similarity to a Src SH3 binding sequence
                                                                                                                  The present invention discloses a polypeptide-guanosine triphosphatase activator protein-10.01. The invention also discloses the method for curing several diseases, such as squamobasal cell carcinoma of skin, osteosarcoma, leukemia and teratoma by using said polypeptide. The present sequence represents the N-terminal sequence of guanosine triphosphate activator protein 10.01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New DNA encoding proteins that bind to Src-homology domain 3 - used to
                                         A polypeptide-guanosine triphosphatase activator protein -10.01 and polynucleotide for coding this polypeptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     - SH3_binding_motif
"corresponds to consensus sequence RPLPXXP"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Proline-rich SH3 binding peptide fuse5B-4.51 from random library.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              inhibit signal transduction, e.g. in cases of T cell activation
                                                                                                                                                                                                                                                         6; Length 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              domain; SH3 domain; oncogene; consensus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    South VJ,
                                                                                                                                                                                                                                                       48.1%; Score 38; DB 70.0%; Pred. No. 36;
                                                                                                                                                                                                                                                                                     1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Cheadle C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RHON ) RHONE POULENC RORER PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
                                                                                       Example 6; Page 21; 33pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                                                                                          AAW05069 standard; peptide; 15 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 1; Col 7; 39pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                     7; Conservative
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/label=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Searfoss GH;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Src homology domain; SH random display library.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /note=
                                                                                                                                                                                                                                                                                                                   4 HSFGSPRPLP 13
                                                                                                                                                                                                                                                                                                                                                HSLGSLOPLP 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (revised)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1996-361965/36.
                                                                                                                                                                                                                                                                     Local Similarity
                                                                                                                                                                                                                          Sequence 15 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    vashchenko YD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19-APR-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19-APR-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       25-MAR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   17-DEC-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                30-JUL-1996.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   French SM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAW05069;
                                                                                                                                                                                                                                                         Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New human K-ras proto-oncogene protein 10.67 polypeptide for treating malignant tumors, hemopathy, development disorder, human immunodeficiency
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention discloses a new kind of polypeptide, human K-rass and a DNA recombination process to produce the polypeptide. The present invention also discloses applying the polypeptide in treating various diseases, such as malignant tumours, haemopathy, development disorder, human immunodeficiency virus (HIV) infection, immunological diseases and various inflammations. The present invention also discloses the antagonist resisting the polypeptide and its treatment effect. The present invention also discloses application of the polynucleotides present invention also discloses application of the polynucleotides recoding human K-ras proto-oncogene protein 10.67, The present sequence represents human K-ras proto-oncogene protein 10.67, N-terminus, used in an ELISA (enzyme-linked immunosorbent assay) experiment
identified within a protein isolated by screening a lambda-lox mouse embryo CDNA library with a Src SHH domain probe. The present sequence is that of one of the positive peptides identified from the fuse 5B random display library. (Updated on 25-MAR-2003 to correct PI field.)
                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            development disorder; human immunodeficiency virus infection; HIV; immunological disease; inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  virus infection, immunological diseases and various inflammations.
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                                                                                                                                                                 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; K-ras 10.67; proto-oncogene; tumour; haemopathy;
                                                                                                                                 Length
                                                                                                                                                                                                                                                                                                                                                                                                                                           Human K-ras 10.67 proto-oncogene protein, N-terminus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 5; Page 19 (disclosure); 33pp; Chinese.
                                                                                                                                 DB 2;
                                                                                                                             Score 37; DB 2
Pred. No. 51;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (BODE-) BODE GENE DEV CO LTD SHANGHAI.
                                                                                                                                                                                                                                                                                                                              ABG72660 standard; peptide; 15 AA.
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0
                                                                                                                               46.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                 Conservative
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Matches 6; Conservative
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                                                                                                                                                                                                     4 HSFGSPRPLP 13
                                                                                                            Query Match
Best Local Similarity
7; Conserve
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                                                                                             Sequence 15 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                       26-FEB-2003
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Gaps

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2; Indels

7 GSPRPLP 13

8

Jaye MC;

GSPOPLP 14

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AAWI1098-WI1124 are peptides that bind to the Src SH3 domain. The SH3 binding peptides are useful in modulating signal transduction pathways at the cellular level (especially protein tyrosine kinase-mediated), modulating oncogenic protein activity, or providing compounds for the development of drugs with the ability to modulate broad classes, as well as specific classes, of proteins involved in signal transduction and also for regulating the processing, trafficking or translation of RNA. Conjugates of the peptides with detectable labels or imaging agents are useful for imaging cells, tissues and organs in which Src or Src-related
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Peptide with binding affinity for Src homology region 3 (SH3) domains of proteins - useful for e.g. modulating signal transduction pathways at the cellular level, esp. protein tyrosine kinase-mediated.
                                                                                                                      Src SH3 domain-binding peptide used in signal transduction modulation.
                                                                                                                                                   Src; SH3; Src homology region 3; binding affinity; oncogenic protein;
protein tyrosine kinase; signal transduction; RNA processing;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Src homology 3 domain; SH3 domain; peptidomimetic; mimetic; modulator;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                      Der CJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Src homology 3 domain binding nonamer peptide #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              44.3%; Score 35; DB 2;
62.5%; Pred. No. 92;
cive 3; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                    Quilliam LA,
                           AAW11111 standard; peptide; 13 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 38; Page 87; 116pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                  Thorn JM,
                                                                                                                                                                                                                                                                                                           95WO-US009382.
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95US-00483555.
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                                                                                         (first entry)
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                                                                                                                                                                                   trafficking; translation.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 proteins are expressed
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TPRPLPLI
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                                                                                       25-JUN-1997
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                                                                                                                                                                                                                                                                                                         24-JUL-1995;
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                                                                                                                                                                                                                                                                           08-FEB-1996
                                                                                                                                                                                                                  Synthetic.
                                                          AAW11111;
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            AAW11111
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RESULT
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The present invention describes spiro[Pyrrolidine-2':3-piperidin-2-ones], their fused ring analogues, and salts of both, all as individual compounds or their libraries. They are peptidominatics, and interact with protein Src homology 3 (SH3) domains, involved in protein-protein interactions in signal transduction. They cause disruption of the pathway in particular, they inhibit the protein tyrosine kinase system involved in cell growth regulation. The cells may be microbial, e.g. pathogenic bacterial or fungal cells responsible for infections, or of transformed cells, e.g. responsible for neoplastic conditions. Specific examples of Src proteins with SH3 domains are those known by the abbreviations p85, Fyn, Lyn, Hck, Syk, Grb2, Gap, STAT, p47-phox, p67-phox, and Brk. The present sequence represents a nonamer peptide used in an example from the
                                                                                                                                                                                                                                                                                                                                                                                                                        spiro[Pyrrolidine-2':3-piperidin-2-ones] and their fused ring analogues -
used, e.g. to modulate signal transduction protein interactions, inhibit
cell growth and for use in treating infections and cancer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cortactin; SH3 domain; binding peptide; Src homology region 3;
tyrosine kinase; immune response; lymphokine; interleukin 1; Nck; Abl;
PLCgamma; p53bp2; Crk; Yes; Grb2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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beta-turn template; spiro[Pyrrolidine-2':3-piperidin-2-ones]; fused ring analogue; signal transduction protein interaction; inhibit cell growth; infection; cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 43.0%; Score 34; DB 2; Length 9; 100.0%; Pred. No. 1.4e+06; Live 0; Mismatches 0; Indels
                                                                                                    Location/Qualifiers
                                                                                                                                  /note= "acetylated"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 12; Page 59; 73pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAW39011 standard; peptide; 15 AA.
                                                                                                                                                                                                                                         98WO-US010861,
                                                                                                                                                                                                                                                                         97US-00864241.
                                                                                                                                                                                                                                                                                                                                                          Witter DJ;
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                                                                                                                                                                                                                                                                                                                       (CADU-) CADUS PHARM CORP.
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                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1999-080804/07.
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Best Local Similarity
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                                                                                                               Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 9 AA;
                                                                                                                                                                                                                                                                         28-MAY-1997;
                                                                                                                                                                                                                                                                                         18-MAY-1998;
                                                                                                                                                                    WO9854208-A1
                                                                                                                                                                                                      03-DEC-1998
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                                                               Synthetic.
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Gaps . 0 WO9730074-A1

95US-00483555.

07-JUN-1995;

(UYNC-) UNIV NORTH CAROLINA

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The present sequence represents a peptide which resembles a Src homology region 3 (SH3) binding peptide. SH3 binding peptides are selected from:

(a) peptides which bind the SH3 domain of Cortactin; (b) peptides which bind the SH3 domain of Cortactin; (b) peptides which bind the SH3 domain of Src; (e) peptides which bind the SH3 domain of Src; (e) peptides which bind the SH3 domain of Crk;

(b) peptides which bind the SH3 domain of Yes; and (i) peptides which bind the amino-terminal SH3 domain of Crk;

(h) peptides which bind the SH3 domain of Yes; and (i) peptides which bind the amino-terminal SH3 domain of Grb. The purified binding peptides to a used in the method to identify inhibitors of their binding to their respective SH3 domains, which could be used to modulate the pharmacological activity of proteins or polypeptide containing the SH3 domain. The peptides can also be used to activate Src or Src-related protein tyrosine kinases, to stimulate the immune response by increasing the production of certain lymphokines, e.g. tumour necrosis factor-alpha and interleukin-1, or to deliver a conjugated molecule to certain
                                                                                                                                                                                                                                                           Src homology region 3 binding peptide - used to activate Src tyrosine kinase(s) and to stimulate immune response by increasing production of certain lymphokine(s), e.g. interleukin-1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Src SH3 domain-binding peptide used in signal transduction modulation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Src; SH3; Src homology region 3; binding affinity; oncogenic protein;
protein tyrosine kinase; signal transduction; RNA processing;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                            Fowlkes DM;
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                                                                                                                                                                           Der CJ,
                                                                                                                                                                         Quilliam LA,
                                                                                                                                                                                                                                                                                                                              Claim 22; Page 93; 131pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAW11105 standard; peptide; 13 AA.
                                                                                                                                                                       Sparks AB, Kay BK, Thorn JM,
                                                                                   96US-00602999
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                                                                                                                                    UYNC-) UNIV NORTH CAROLINA
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Best Local Similarity 100...
6; Conservative
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                                                                                                                    CYTO-) CYTOGEN CORP
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                                                 14-FEB-1997;
                                                                                   16-FEB-1996;
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                21-AUG-1997
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                                                                                                                 Peptide with binding affinity for Src homology region 3 (SH3) domains of proteins - useful for e.g. modulating signal transduction pathways at the cellular level, esp. protein tyrosine kinase-mediated.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Peptide with binding affinity for Src homology region 3 (SH3) domains of proteins - useful for e.g. modulating signal transduction pathways at the cellular level, esp. protein tyrosine kinase-mediated.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Src SH3 domain-binding peptide used in signal transduction modulation.
                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Src; SH3; Src homology region 3; binding affinity; oncogenic protein;
protein tyrosine kinase; signal transduction; RNA processing;
                                                                                                                                                                                                                                                                                                                                                                                                                                        .,
                                                                                                                                                                                                                                                                                                                                                                                      41.8%; Score 33; DB 2; Length 13; 62.5%; Pred. No. 1.9e+02;
                                                           Der CJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Der CJ;
                                                           Quilliam LA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Quilliam LA,
                                                                                                                                                                             Claim 38; Page 85; 116pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAW11098 standard; peptide; 14 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 32; Page 79; 116pp; English.
                                                           Thorn JM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Thorn JM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         94US-00278865.
95US-00483555.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               95WO-US009382
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (UYNC-) UNIV NORTH CAROLINA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 25-JUN-1997 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                    5; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       trafficking; translation.
                                                                                                                                                                                                                                                                                                                                            proteins are expressed
                                                         Kay BK,
                                                                                    WPI; 1996-117151/12.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                8 SPRPLPW 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2 TPRPLPML 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Kay BK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1996-117151/12.
                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                         Sequence 13 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            24-JUL-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         22-JUL-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       07-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO9603649-A1
                                                         Sparks AB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sparks AB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
à
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AAWI1098-WI1124 are peptides that bind to the Src SH3 domain. The SH3 binding peptides are useful in modulating signal transduction pathways at the cellular level (especially protein tyrosine kinase-mediated), modulating oncogenic protein activity, or providing compounds for the development of drugs with the ability to modulate broad classes, as well as specific classes, of proteins involved in signal transduction and also for regulating the processing, trafficking or translation of RNA. Conjugates of the peptides with detectable labels or imaging agents are useful for imaging cells, tissues and organs in which Src or Src-related
                                                                                                                                                                                                                                                                                    proteins are expressed
            889999999998888
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Sequence 14 AA;

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41.8%; Score 33; DB 2; Length 14; 62.5%; Pred. No. 2.1e+02; ive 3; Mismatches 0; Indels
: | | | | | | ::
3 TPRPLPML 10
                                                8 SPRPLPW 15
                                                 à
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0;

Gaps

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RESULT 12

AAW25512 standard; peptide; 14 AA. AAW25512

AAW25512;

SH3 synthetic binding peptide. (first entry) 27-MAR-1998

Cortactin; SH3 domain; binding peptide; Src homology region 3; tyrosine kinase; immune response; lymphokine; interleukin 1; Nck; Abl; PLCgamma; p53bp2; Crk; Yes; Grb2.

Synthetic.

WO9730074-A1.

21-AUG-1997.

97WO-US002298. 14-FEB-1997;

96US-00602999 16-FEB-1996;

UNIV NORTH CAROLINA (CYTO-) CYTOGEN CORP. (DXNC-)

Quilliam LA, Der CJ, Fowlkes DM; Sparks AB, Kay BK, Thorn JM, Rider JE;

WPI; 1997-424972/39.

Src homology region 3 binding peptide - used to activate Src tyrosine kinase(s) and to stimulate immune response by increasing production of certain lymphokine(s), e.g. interleukin-1. Src homology

Disclosure; Fig 7; 131pp; English.

(Str. homology region 3) binding peptides are selected from: (a) peptides which bind the SH3 domain of Cortactin, (b) peptides which bind the Mal domain of Cortactin, (b) peptides which bind the Mal domain of McK; (c) peptides which bind the SH3 domain of Ab1, (d) peptides which bind the SH3 domain of Str. (e) peptides which bind the SH3 domain of Crk; (e) peptides which bind the Mal domain of Crk; (i) peptides which bind the MH3 domain of Crk; (ii) peptides which bind the MH3 domain of Crk; (ii) peptides which bind the SH3 domain of Grb2. The purified binding peptides can the amino-terminal SH3 domain of Grb2. The purified binding peptides can be used in the method to identify inhibitors of their binding to their pharmacological activity of proteins or polypeptide containing the SH3 domain. The peptides can also be used to activate Src or Src-related the present sequence represents a Src SH3 synthetic binding peptide. SH3

polypeptides in samples

Sequence 14 AA;

```
AA173060 to AA179867 represent isolated human polymorphic polymuclectide sequences [1], which contain single nuclectide polymorphisms (SNPs).

AAW53114 to AAM53329 represent peptides related to human polymorphic polymorphic polymorphic polymorphic polymorphic sequences. The sequences can be used in protein the prevention. [1] and the polypeptides encoded by therapy, and in vaccine production. [1] and the polypeptides encoded by expensive the properties of diseases associated with inappropriate expression of polymorphic polypeptides. For example, [1] may be used to treat disorders by rectifying mutations or deletions in a patient's genome that affect the activity of polypeptides by expressing inactive proteins or to supplement the patients own production of polypeptides. Additionally, [1] and its complementary and quantitate the presence of similar nucleic assays to detect therefore which patients may be in need of restorative therapy. The preference may be used as antigens in the production of polypeptides encoded by [1] may be used as antigens in the production of antithedian encoded by [1] may be used as antigens in the production of antithedian encoded by [1] may be used as antigens in the production of
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protein tyrosine kinases, to stimulate the immune response by increasing the production of certain lymphokines, e.g. tumour necrosis factor-alpha and interleukin-1, or to deliver a conjugated molecule to certain cellular compartments containing Src or Src related proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human nonconservative amino acid changing SNP related peptide SEQ:6990.
                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human, single nucleotide polymorphism; SNP; genome, gene therapy, protein therapy; vaccine; probe; diagnostic assay; detection; quantitation; restorative therapy; polymorphic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Polymorphic nucleic acid sequences, useful in genetic testing and
                                                                                                                                                                                          .
                                                                                                                                               Score 33; DB 2; Length 14;
Pred. No. 2.1e+02;
3; Mismatches 0; Indels
                                                                                                                                                 DB 2; Length 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 29; Page 2643; 2653pp; English.
                                                                                                                                                                                                                                                                                                                                                                            AAM53295 standard; peptide; 14 AA.
                                                                                                                                                 41.8%;
62.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       30-NOV-2000; 2000WO-US032758.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 29-NOV-2000; 2000US-00726173.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                99US-0168138P
                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                        5; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                             8 SPRPLPVV 15
                                                                                                                                                                                                                                                                      3 TPRPLPML 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2001-356160/37.
                                                                                                                                                                  Best Local Similarity
                                                                                                        Sequence 14 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200140521-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              30-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            09-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             07-JUN-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                   AAM53295;
                                                                                                                                               Query Match
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                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                 RESULT 13
                                                                                                                                                                                                                                                                                                                                                      AAM53295
  8886688
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Gaps

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Treating a cancer patient involves administering to patient a cytotoxic T-lymphocyte, an antigen presenting cell that activates T cells, a peptide of cytochrome P450 lBl, or a polynucleotide encoding the peptide.
                                                                                                                                                                                                        Cytochrome P450 1B1; CYP1B1; cancer; immunotherapy; diagnosis; HLA; human leukocyte antigen; cytochoxic T-lymphocyte; CTL; MHC; APC; major histocompatibility complex; antigen presenting cell; melanoma; lymphoma; carcinoma; sarcoma; multiple myeloma; leukaemia.
                                                                                                                                                                                                                                                                                                                                                                                                  Maecker B;
 41.8%; Score 33; DB 4; Length 14;
85.7%; Pred. No. 2.16+02;
"Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                 Sherr D, Nadler LM,
                                                                                                                                                                                     CYP1B1 HLA-A2 binding peptide fragment #35.
                                                                                                                                                                                                                                                                                                                                                              (DAND ) DANA FARBER CANCER INST INC.
                                                                                                                        AAM06985 standard; peptide; 10 AA.
                                                                                                                                                                                                                                                                                                                                                                                             Schultze JL, Vonderheide RH,
Von Bergwelt- Baildon M;
                                                                                                                                                                                                                                                                                                                       15-NOV-2000; 2000WO-US031513.
                                                                                                                                                                                                                                                                                                                                            99US-0165590P
Overy Match
Best Local Similarity 85...
6; Conservative
                                                                                                                                                                 (first entry)
                                                   8 SPRPLPV 14
                                                                       6 SPRPAPV 12
                                                                                                                                                                                                                                                                                                                                                                         (UYBO-) UNIV BOSTON
                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2001-355537/37
                                                                                                                                                                                                                                                                               WO200135810-A2.
                                                                                                                                                                                                                                                                                                                                          15-NOV-1999;
                                                                                                                                                                                                                                                            Homo sapiens.
                                                                                                                                                                 09-OCT-2001
                                                                                                                                                                                                                                                                                                   25-MAY-2001.
                                                                                                                                             AAM06985;
                                                                                                                AAM06985
                                                   à
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The present invention describes a method for treating a patient having, or is at risk of having, a cell expressing cytochrome P450 1B1 (CYPIB1), which involves adminiatering to the patient a cytocoxic T-lymphocyte (CTL) that kills the cell, an antigen presenting cell (APC) that cell, a peptide of CYPIB1 that binds to major histocompatibility activates CTL, a peptide of CYPIB1 that binds to major histocompatibility complex (MHC) molecule or CYPIB1 or peptide of CYPIB1. Also described are: (1) a method for assessing the complex molecule, involving measuring the level of CYPIB1 or the peptide of CYPIB1 or a peptide of CYPIB1 that binds to MHC complex molecule, involving measuring the level of CTL cspecific for CYPIB1 or the peptide of CYPIB1 in a sample from the patient cspecific of CYPIB1 or the specifically kills a cell expressing CYPIB1 or a specific of a MHC complex molecule. The method is useful for treating a patient or a specific of a MHC complex-restricted fashion; and (4) and extinct complex molecule. The method is useful for treating a patient complex molecule. The method is useful for treating a patient complex molecule. The method is useful for treating a patient of a useful for the prevention, treatment and diagnosis of cancer. Graving or is at risk of having a cell that expresses CYPIB1. The method is useful for the prevention, treatment and diagnosis of cancer. Graving or arian, uterime, cervical, prostate, liver, colon, pancreatic and clung, ovarian, uterime, cervical, prostate, liver, colon, pancreatic and colon, pancreatic and chain cancer. AAM05005 to AAM13566 represent CYPIB1 peptides which can be an order of the present invention and context of sequence used in the exemplification of the present invention

Disclosure; Page 40; 121pp; English.

Sequence 10 AA;

Sequence 10 AA;

```
The present invention describes a method for treating a patient having, or is at risk of having, a cell expressing cytochrome P450 IBI (CYPIBI), which involves administering to the patient a cytocoxic T-lymphocyte (CTL) that kills the cell, an antigen presenting cell (APC) that complex (MTC) molecule, or a nucleic acid molecule encoding CYPIBI or complex (MMC) molecule, or a nucleic acid molecule encoding CYPIBI or peptide of CYPIBI. Also described are: (1) a method for assessing the level of immunity of a patient to CYPIBI or a peptide of CYPIBI that complex molecule, involving measuring the level of CYPIBI or the peptide of CYPIBI in a sample from the patient of peptide of CYPIBI or the peptide of CYPIBI in a sample from the patient of complex molecule (1) that binds to MHC complex molecule; (3) an exvivation a specific of a MHC complex restricted fashion; and (4) and exvivation as specific of a MHC complex molecule. The method is useful for treating a patient can any or is at risk of having a cell that expresses CYPIBI. The method is useful for the prevention, treatment and diagnosis of cancer, e.g. melanoma, lymphoma, carcinoma, sarcoma, multiple myeloma, leukemia, and cludy ovarian, uterine, cervical, prostate, liver, colon, panorcatic and brain cancer. AAMO6505 to AAMI3566 represent CYPIBI peptides which can brain cancer and in the exemplification of the present invention
                                                              0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Treating a cancer patient involves administering to patient a cytotoxic T ~1ymphocyte, an antigen presenting cell that activates T cells, a peptide of cytochrome P450 1B1, or a polynucleotide encoding the peptide.
                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                           Cytochrome P450 1B1; CYP1B1; cancer; immunotherapy; diagnosis; HLA; human leukocyte antigen; cytotoxic T-lymphocyte; CTL; MHC; APC; major histocompatibility complex; antigen presenting cell; melanoma; lymphoma; carcinoma; sarcoma; multiple myeloma; leukaemia.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Maecker
                         Length 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sherr D, Nadler LM,
                       Score 32; DB 4; I
Pred. No. 2.1e+02;
0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (DAND ) DANA FARBER CANCER INST INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Page 69; 121pp; English.
                                                                                                                                                                                                                              AAM08985 standard; peptide; 10 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Vonderheide RH,
                   40.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      15-NOV-2000; 2000WO-US031513.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               99US-0165590P.
Query Match
Best Local Similarity 70.0%,
Ass 7; Conservative
                                                                                                                                                                                                                                                                                                     09-OCT-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Von Bergwelt- Baildon M;
                                                                                                                                                                                                                                                                                                                                          HLA-A *0201 decamer #2.
                                                                                                15
                                                                                                                                   1 FLDPRPLTVV 10
                                                                                            6 FGSPRPLPVV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (UYBO-) UNIV BOSTON.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2001-355537/37.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO200135810-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Schultze JL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  25-MAY-2001.
                                                                                                                                                                                                                                                                   AAM08985;
                                                                                                                                                                                          RESULT 15
                                                                                                                                                                                                              AAM08985
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0; Gaps Ouery Match 40.5%; Score 32; DB 4; Length 10; Best Local Similarity 70.0%; Pred. No. 2.1e+02; Matches 7; Conservative 0; Mismatches 3; Indels

0;

6 FGSPRPLPVV 15 | | | | | | | 1 FLDPRPLTVV 10

qq δ

Search completed: April 2, 2004, 10:08:42 Job time: 47 secs

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GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
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OM protein - protein search, using sw model

Run on:

(without alignments) 131.170 Million cell updates/sec April 2, 2004, 10:05:03; Search time 11 Seconds

US-10-009-049-2

1 SRAHSFGSPRPLPVV 15 Perfect score: Title:

BLOSUM62 Sequence:

Gapop 10.0 , Gapext 0.5 Scoring table:

283366 seqs, 96191526 residues Searched:

Total number of hits satisfying chosen parameters:

2522

Minimum DB seq length: 0 Maximum DB seq length: 15

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

1: pir1:*
2: pir2:*
3: pir3:*
4: pir4:* PIR 78:* Database

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

	Description	hypothetical prote	O	protein OF200053 -	bradykinin-potenti	bradykinin-potenti	kinesin liqht chai	photosystem II oxy	Î-cell receptor be	protein 0F200037 -	1.3	orphin (ਨ	Ig kappa chain V-I	photosystem II oxy	deoxynúcleotidyltř	Ig heavy chain CRD	locustamyotropin 1	platelet-derived q	dermorphin - Rohde	leucokinin VI - Ma	orf 1 rara 5'-regi	acylglyc	173K exoantigen -	photosystem II 3.7	I-cell receptor be	receptor	a i	formate dehydrogen	orf19 3' of eryK -
	Д	S27140	XAVI9B	PA0089	C37196	D37196	S43170	PA0013	PH1457	PA0060	PS0452	S36662	D47393	PL0157	G44957	184603	PT0238	B61620	A22789	A61324	JS0316	S15594	PX0030	I33098	S01122	PH1459	PH1468	PC2121	S59492	A40634
	DB	2	Н	7	7	7	7	~	N	N	N	(1	N	N	~	7	~	7	7	-	~	4	7	7	C4	7	7	4	7	۲,
	Query Match Length	14	13	13	11	11	12	14	12	15	15	7	80	13	14	13	6	13	15	7	80	9	10	11	12	12	12	12	15	15
₩,	Query Match	34.2	32.9	32.9	30.4	30.4	30.4	30.4	29.1	ο.	28.5	7.	7.	7.	7.	7	ė	26.6	ç.	ű.	25.3	ď.	25.3	'n.	25.3	'n	2		25.3	
	Score		26		24	24	24	24	23	23	22.5	22	22	22	N	21.5	21	21	21	20	20	20	20	20	20	20	20	20	20	20
;	Result No.	! !	7	m	4	വ	ø	7	œ	6	10	11	12	13			16	17		19	20	21	22	23					28	

hypothetical prote	mixed lymphocyte r	xylose isomerase (angiotensin-conver	bradykinin-potenti	substance P - chic	substance P - rain	substance P - Atla	probable antigen 5	probable substance	T-cell receptor be	T-cell receptor be	Iq lambda chain J	T-cell receptor be		T-cell receptor be
B32800	PX0031	A31576	XAVI6B	XASNBA	JN0023	S23308	\$23306	E41476	833300	S26559	S26554	A37933	PH1467	PH1462	PH1469
7	~	Ŋ	7	Н	N	7	7	7	7	7	7	7	N	7	~
15	15	9	10	11	11	11	11	11	11	12	12	12	12	12	12
25.3	25.3	24.1	24.1	24.1	24.1	24.1	24.1	24.1	24.1	24.1	24.1	24.1	24.1	24.1	24.1
25	•														
20 25		19	19	19	19	19	19	19	19	19	19	19	19	19	19

ALIGNMENTS

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R,Keaveney, M.; Klug, J.; Gannon, F.
DNA Seq. 2, 347-358, 1992
NA Seq. 2, Sequence analysis of the 5' flanking region of the human estrogen receptor gene.
A;Reference number: S27140; WUID:93075998; PMID:1476547
A;Accession: S27140
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ;
                            C;Species: Homo sapiens (man)
C;Date: 05-Mar-1994 #sequence_revision 12-May-1995 #text_change 18-Aug-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                 A;Cross-references: EMBL:X62462; NID:931201; PIDN:CAA44319.1; PID:931202
C;Superfamily: unassigned leader peptides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               .,
                                                                                                                                                                                                                                                                                                                                                                                                                     34.2%; Score 27; DB 2; Length 14; 100.0%; Pred. No. 2.7e+02; tive 0; Mismatches 0; Indels
hypothetical protein 1 estrogen receptor 5'-region - human
                                                                                                                                                                                                                                                A;Status: translation not shown
A;Molecule type: DNA
A;Residues: 1-14 <KEA>
                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity 100.0
Matches 5; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2 RAHSF 6
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                                                                                   C; Accession: S27140
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RESULT 2

angiotensin-converting enzyme inhibitor V-9 - jararaca C;Species: Bothrops jararaca (jararaca) C;Date: 13-Jul-1981 #sequence_revision 13-Jul-1981 #text_change 08-Dec-1995

Cincession: A01253
Riondetti, M.A.; Williams, N.J.; Sabo, E.F.; Pluscec, J.; Weaver, E.R.; Kocy, O.
Biochemistry 10, 4013-4039, 1971
Biochemistry 10, 4013-4039, 1971
Biochemistry 10, 4013-4039, 1971
Biochemistry 10, 4013-4039, 1971
Biochemistry 10, 4013-403
Birtherance number: A90356; MUID:72118526; PMID:4334402
Birtherance number: A90356; PMID:434402
Birtherance number: A90356; PMID:434402
Birtherance number: A90356; PMID:72118526; PMID:72118526; PMID:72118526; PMID:72118526; PMID

.. 0 h 32.9%; Score 26; DB 1; Length 13; Similarity 55.6%; Pred. No. 3.6e+02; 5; Conservative 1; Mismatches 3; Indels Best Local Similarity Matches 5; Conserv Query Match

o O

Gaps

7 GSPRPLPW 15 3 GWPRPGPEI 11 δ d

RESULT 3

Query Match 32.9%; Score 26; DB 2; Length 13; Best Local Similarity 80.0%; Pred. No. 3.6e+02; Matches 4; Conservative 1; Mismatches 0; Indels

7 GSPRP 11

5 GAPRP 9

A; Molecule type: protein A; Residues: 1-13 <CHO>

```
R;Chernajovsky, Y.; Brown, A.; Jones, T.A.; Sheer, D. submitted to the EMBL Data Library, December 1992
A;Description: Promoter first exon/intron characterization and chromosomal location of the A;Reference number: 843170
A;Accession: 843170
A;Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         R;Kamo, M.; Kawakami, T.; Miyatake, N.; Tsugita, A. submitted to JIPID, July 1994
A;Description: Separation and characterization of Arabidopsis proteins by two-dimensional
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           T-cell receptor beta chain (clone A3/IC1) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 10-Mar-1994 #sequence_revision 10-Mar-1994 #text_change 17-Mar-1999
C;Accession: PH1457; S26542
R;Casanova, J.L.; Martinon, F.; Gournier, H.; Barra, C.; Pannetier, C.; Regnault, A.; Kon J. Exp. Med. 177, 811-820, 1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A, Experimental source: cytolytic T-lymphocyte, clone A3/IC1
R; Casanova, J.L.; Cerottini, J.C.; Matthes, M.; Necker, A.; Gournier, H.; Barra, C.; Widn
J. Exp. Med. 176, 439-447, 1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A,Title: T cell receptor selection by and recognition of two class I major histocompatibi A,Reference number: PH1430, MUID:93171821, PMID:8436911
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A,Reference number: S26512, MUID:92364546, PMID:1380061
A,Accession: S26542
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C;Date: 13-Jan-1995 #sequence_revision 18-Aug-2000 #text_change 18-Aug-2000
C;Accession: S43170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       C;Species: Arabidopsis thaliana (mouse-ear cress)
C;Date: 30-Jun-1992 #sequence_revision 06-Jan-1995 #text_change 23-Mar-1995
C;Accession: PA0013
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A; Cross-references: EMBL: X69658; NID: 9468786; PIDN: CAA49349.1; PID: 9468787
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A;Residues: 1-14 «KAM»
A;Experimental source: leaf
C;Keywords: photosynthesis; photosystem II
                                                                                                                                                                      kinesin light chain - human (fragment)
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A; Residues: 1-12 <CAS>
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3 GPPRP
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Richow, L. P.; Fukaya, N.; Sugiura, Y.; Ueno, Y.; Tabuchi, K.; Tsugita, A.

Submitted to JTPID, October 1994

A; Description: Two dimensional polyacrylamide gel electropheresis of Fusarium sporotrich

A; Reference number: PA0051

A; Accession: PA0089
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C;Species: Bothrops insularis (island jararaca)
C;Date: 14-Feb-1992 #sequence_revision 01-Dec-1992 #text_change 05-Aug-1994
C;Date: 14-Feb-1992 #sequence_revision 01-Dec-1992 #text_change 05-Aug-1994
C;Cacession: D37196
B;Cintra, A.C.O.; Vieira, C.A.; Giglio, J.R.
J. Protein Chem. 9, 221-227, 1990
A;Pitle: Primary structure and biological activity of bradykinin potentiating peptides A;Pitle: Primary structure and biological activity of bradykinin potentiating peptides A;Reference number: A37196; MUID:90351557; PMID:2386615
A;Reference number: A37196; MUID:90351557; PMID:2386615
A;Accession: D37196
A;Section: proliminary
A;Residues: 1-11 cIIN>
C;Keywords: pyroglutamic acid
F;1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental
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0
                                                                                            protein QF200053 - fungus (Fusarium sporotrichioides) (fragment)
C;Species: Fusarium sporotrichioides
C;Date: 20-Feb-1995 #sequence_revision 20-Feb-1995 #text_change 23-Mar-2001
C;Accession: PA0089
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n 30.4%; Score 24; DB 2; Length 11; Similarity 80.0%; Pred. No. 6.5e+02; 4; Conservative 0; Mismatches 1; Indels

Best Local Similarity Matches 4; Conserv

Query Match

7 GSPRP 11

3 GPPRP 7

A)Status: preliminary
A;Molecule type: protein
A;Residues: 1-11 <CIN>
C:Keywords: pyroglutamic acid
F;I/Modified site: pyrrolidone

bradykinin-potentiating peptide 4 - island jararaca

Ouery Match
30.4%; Score 24; DB 2; Length 11;
Best Local Similarity 80.0%; Pred. No. 6.5e+02;
Matches 4; Conservative 0; Mismatches 1; Indels

7 GSPRP 11

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4 HSFGSPR 10
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1 YAFGYPK 7
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G44957
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                                                                                                                                                                                                                                                                                                                                                                                                   protein QF200037 - fungus (Fusarium sporotrichioides) (fragment)
C;Species: Fusarium sporotrichioides
C;Species: Co-Feb-1995 #sequence_revision 20-Feb-1995 #text_change 23-Mar-2001
C;Accession: PA0060
R;Chow, L.P.; Fukaya, N.; Sugiura, Y.; Ueno, Y.; Tabuchi, K.; Tsugita, A.
Submitted to JIPID, October 1994
A;Description: Two dimensional polyacrylamide gel electropheresis of Fusarium sporotrich
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        dermorphin (Lys-7) (validated) - two-colored leaf frog
C;Species: Phyllomedusa bicolor (two-colored leaf frog)
C;Date: 10-Dec-1993 #sequence_revision 13-Mar-1997 #text_change 18-Aug-2000
C;Accession. S3666
B;Mignogna, G:; Severini, C.; Simmaco, M.; Negri, L.; Falconieri Erspamer, G.; Kreil, G.
FEBS Lett. 302, 151-154, 1992
A;Title: Identification and characterization of two dermorphins from skin extracts of th
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C;Species: Oryza sativa (rice)
C;Date: 0.3-Feb-1994 #sequence_revision 03-Feb-1994 #text_change 11-Apr-1995
C;Accession: PS0452
R;Tsugita, A, Mygatake, N.
Submitted to JIPDD, April 1993
A;Reference number: PS0208
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 15;
                                                                                                                                                                  29.1%; Score 23; DB 2; Length 12; 83.3%; Pred. No. 1e+03;
                                                                              A; Experimental source: cytolytic T-lymphocyte, clone Cw3/Cas2 C; Superfamily: immunoglobulin homology C; Keywords: receptor; T-cell
                                                                                                                                                                                                                 1; Indels
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Pred. No. 1.3e+03;
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100.0%; Pred. No. 1....
... 0; Mismatches
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A,Molecule type: protein
A,Residues: 1-15 CTS0
C,Comment: molecular weight 32K, pI 5.3.
                                                        A; Cross-references: EMBL:X67992
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Best Local Similarity 100.v
                                                                                                                                                                                                             5; Conservative
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A; Accession: PA0060
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Matches 5; Conserv
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             A; Molecule type: mRNA
A; Residues: 1-12 < CA2>
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Status: preliminary
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R;Duve, H.; Johnsen, A.H.; Scott, A.G.; Yu, C.G.; Yagi, K.J.; Tobe, S.S.; Thorpe, A. Proc. Natl. Acad. Sci. U.S.A. 90, 2456-2460, 1993
A;Title: Callatostatins: neuropeptides from the blowfly Calliphora vomitoria with sequent A;Reference number: A47393; MUID:93211980; PMID:8460157
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      "g kappa chain V-II region (anti-myelin-associated glycoprotein, PEC) - human (fragment) C;Species: Homo sapiens (man)
C;Species: Homo sapiens (man)
C;Accession: PLO157; C61458
R;Brouet, J.C.; Dellagi, K.; Gendron, M.C.; Chevalier, A.; Schmitt, C.; Mihaesco, E.
J. Exp. Med. 170, 1551-1558, 1989
A;Title: Expression of a public idiotype by human monoclonal IgM directed to myelin-association number: A61458; MUID:90039128; PMID:2478651
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A;Molecule type: protein
A;Residues: 1-13 <BR2>
C;Comment: This protein is one of monoclonal IgM reactive with myelin-associated glycopr
                            A, Accession: $36662
A, Status: preliminary
A, Molecule type: protein
A, Residues: 1-7 < MIG>
C, Superfamily: dermorphin precursor; dermorphin precursor amino-terminal homology
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       C;Species: Calliphora vomitoria
C;Date: 16-Feb-1994 #sequence_revision 18-Nov-1994 #text_change 28-Apr-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Molecule type: protein
A;Residues: 1-8 <DUV>
A;Experimental source: thoracic ganglia
A;Note: sequence extracted from NCBI backbone (NCBIP:128479)
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A; Reference number: S21152; MUID: 92339502; PMID: 1633846
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Mismatches
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Best Local Similarity 66./~
'--a 4, Conservative
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photosystem II oxygen-evolving complex protein 2 - common tobacco (cv. Samsun NN) (fragm Cispeciaes: Nicotiana tabacum (common tobacco)
C;Date: 03-Jun-1993 #sequence_revision 03-Jun-1993 #text_change 18-Jun-1993
C;Accession: 644957
R;Takeda, S; Sato, F; Ida, K.; Yamada, Y.
R;Takeda, S.; Sato, F; Ida, K.; Yamada, Y.
A;Title: Characterization of polypeptides that accumulate in cultured Nicotiana tabacum A;Reference number: A44957
A;Accession: 644957
A;Accession: G44957
A;Accession: G44957
A;Residues: 1-14 c7AK>

ö 0, 27.8%; Score 22; DB 2; Length 14; 57.1%; Pred. No. 1.8e+03; ative 1; Mismatches 2; Indels Query Match
Best Local Similarity 57.1;
Matches 4; Conservative

3 AHSFGSP 9

6 ANVFGKP 12

Db à

I84603

Georganic leotidyltransferase - human (fragment)
C.Species: Homo sapiens (man)
C.Species: Homo sapiens (man)
C.Species: Homo sapiens (man)
C.Accession: 184603
R.Koiwai, O.; Kaneda, T.; Morishita, R.
Biochem. Biophys: Res. Commun. 144, 185-190, 1987
A.;Title: Analysis of human terminal deoxynucleotidyl transferase cDNA expressible in mam A.;Title: Analysis of human terminal deoxynucleotidyl transferase cDNA expressible in mam A.;Title: Analysis of human terminal deoxynucleotidyl transferase cDNA expressible in mam A.;Title: Analysis of human terminal deoxynucleotidyl transferase CDNA expressible in mam A.;Title: Analysis of human terminal deoxynucleotidyl transferase CDNA expressible in mam A.;Title: Analysis of human terminal deoxynucleotidyl transferase CDNA expressible in mam A.;Cacesic in 1942
A.;Cacetics: Translated from GB/EMBL/DDBJ
A.;Coss-references: GBB:119100; OMIM:187410
A.;Coss-references: GBB:119100; OMIM:187410
A.;Morposition: 10q23-10q24

27.2%; Score 21.5; DB 2; Length 13; 45.5%; Pred. No. 2e+03; tive 2; Mismatches 1; Indels 3; Gaps Query Match Best Local Similarity 45.5 Matches 5; Conservative

7 GSP---RPLPV 14

2 GOPDGROPLPM 12

Search completed: April 2, 2004, 10:11:51 Job time: 11 secs

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GenCore version 5.1.6
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OM protein - protein search, using sw model

April 2, 2004, 09:58:38; Search time 7.66667 Seconds (without alignments) 101.876 Million cell updates/sec Run on:

US-10-009-049-2 79 1 SRAHSFGSPRPLPVV 15 Title: Perfect score:

Sequence:

Scoring table:

141681 seqs, 52070155 residues BLOSUM62 Gapop 10.0 , Gapext 0.5 Searched:

801 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 15

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

SwissProt_42:*

Database :

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

		ď			SUMMARIES		
Result No.	Score	Query Match	Length	DB	ID	Description	ion
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11	23	an.	12	Н	PSP3 PHYPA	a	hyscomitre
12	22	_	60	Н	ALL4_CALVO	0	calliphora
13	22	27.8	10	Н	FAR6_PANRE	_	panaqrellus
14	21	ະດ	11	Н		=#	hoplobatrac
15	21	เก	12	ч	PPK4_PERFU	_	periplaneta
16	21	lo.	12	Н	TIN2_HOPTI		hoplobatrac
17	21	lo.	12		TIN3_HOPTI		hoplobatrac
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58	19	**	80	П	AL18_CARMA		
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30	13	-**	0	-	XYLA_STRSQ	P19149 B	streptomyce
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ALIGNMENTS

O STANDARD; PRT; 8 AA.	000 (Rel. 39, Created) 000 (Rel. 39, Last sequence update) 000 (Rel. 39, Last annotation update)	Cydlastatin 3. Cydlastatin 3. Eukaryota, peranoella (Codling moth). Eukaryota, Metazoa, Arthropoda, Hexapoda, Insecta, Pterygota; Neoptera, Endopterygota; Lepidoptera, Glossata, Ditrysia, Tortricoidea; Tortricidae, Olethreutinae; Cydia. NUEL TaxID=82600;	TISSUE-Larva; TISSUE-Larva; MEDLINE=98054539; PubMed=9392829; MEDLINE=98054539; PubMed=9392829; Dave H., Johnsen A.H., Maestro JL., Scott A.G., Winstanley D., Davey M., East P.D., Thorpe A.; "Lepidoptran peptides of the allatostatin superfamily."; Feptides 18:1301-1309(1997) SIMILARITY: Belongs to the allatostatin family. Neuropeptide; Amidation. Neuropeptide; Amidation. AMIDATION. SEQÜENCE 8 AA; 926 MW; C92879D5AB477415 CRC64;	32.9%; Score 26; DB 1; Length 8; Similarity 71.4%; Pred. No. 1.4e+05; 5; Conservative 1; Mismatches 1; Indels 0; Gaps	SRAHSFG 7 : SRPYSFG 7	BOTJA BPP1 BOTJA STANDARD; PRT; 13 AA. BPP1 BOTJA BPP1 BOTJA STANDARD; PRT; 13 AA. 21-UUL-1986 (Rel. 01, Created) 01-FEB-1994 (Rel. 28, Last sequence update) 28-FEB-2003 (Rel. 41, Last annotation update) 28-FEB-2003 (Rel. 41, Last annotation update) Bradykinin-potentiating peptide S3,1 (13A) (Angiotensin-converting enzyme inhibitor V-9). Bothrops jararaca (Jararaca), and Bothrops jararaca (Jararaca), and Bothrops jararaca (Jararaca). Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Lepdosauria; Squamata; Scheroglossa; Serpentes; Colubroidea; Viperidae; Crotalines; Bothrops. Viperidae; Crotalines; Bothrops. SEGUENCE. SEGUENCE. SEGUENCE. SEGUENCE. MEDLINE=71118526; PubMed=4334402; Ondetti M.A., Williams N.J., Sabo E.F., Pluscec J., Weaver E.R., Kocy O.;
TT 1 CYDPO ALL3 CYDPO	30-MAY-2000 30-MAY-2000 30-MAY-2000 30-MAY-2000	Cydiastatin Cydia pomone Eukaryota; M Neoptera; En Tortricoidea NCBI TaxID=8	SEQUENCE. TISSUE_Larva; MEDLINE=98054539; Duve H., Johnsen Davey M., East P. "Lepidopteran pep- Peptides 18:1301- '- SIMILARITY: B NOUNCOPEPTIDE; Ami- MOD_RES SEQUENCE 8 AA;	Query Match Best Local Simi Matches 5;	1 SRA 	
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Pernollet J.-C., Zivy M., de Vienne D.;
"The maize two dimensional gel protein database: towards an integrated
                                               "Angiotensin-converting enzyme inhibitors from the venom of Bothrops jararaca. Isolation, elucidation of structure, and synthesis."; Biochemistry 10:4033-4039(1971).
                                                                                                                                                                                           Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                  Theor. Appl. Genet. 93:997-1005(1996).
-!- MISCELANEOUS: On the 2D-gel the determined pl of this unknown protein is: 4.9, its Mw is: 37.6 kDa.
Maize-2DPAGE; P80635; COLEOPTILE.
                                                                                                                                                                                                                                                                                                                                      Bukaryota, Viridiplantae, Streptophyta, Embryophyta, Tracheophyta, Peprantophyta, Pagermatophyta, Magnoliophyta, Liliopsida, Poales, Poaceae, PAGCAD clade, Panicoideae, Andropogoneae, Zea.
WCBI_TaxID=4577;
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0
                                                                                                                                                                                                                                                                                   01-OCT-1996 (Rel. 34, Created)
01-OCT-1996 (Rel. 34, Last sequence update)
15-WAR-2004 (Rel. 43, Last annotation update)
Unknown protein from 2D-page of etiolated coleoptile (Spot 45)
                                                                                                                                                                       Score 26; DB 1; Length 13;
Pred. No. 1.9e+02;
1; Mismatches 3; Indels
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Pred. No. 2.2e+02;
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                                           SPECIES=B.insularis; TISSUE=Venom;
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55.6%;
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                                                                                                                                                                                                                                                                                                                                                                                                                          genome analysis program.";
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                                                                                                                                                                                                                                                                    STANDARD;
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1 NPNPVPI 7
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P81668;
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PSBP_PINPS
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MEDLINE=90351557; PubMed=2386615;
Cintra A.C.O., Vieira C.A., Giglio J.R.;
Primary Structure and biological activity of bradykinin potentiating peptides from Bothrops insularis snake venom.";
J. Protein Chem. 9:221-227(1990).
-!- FUNCTION: This peptide both inhibits the activity of the angiotensin-converting enzyme and enhances the action of bradykinin by inhibiting the kinases that inactivate it.

It acts as an indirect hypotensive agent.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Electrophoresis 20:1098-1108(1999).

-!- FUNCTION: May be involved in the regulation of photosystem II.

-!- SUBCELLULAR LOCATION: Chloroplast thylakoid membrane; associated with the photosystem II complex (By similarity).

-!- MISCELLANBOUS: On the 2D-gel the determined pI of this protein (spot NI79) is: 5.9, its MW is: 22 kDa.

-!- SIMILARITY: Belongs to the psbP family.

Photosynthesis; Photosystem II; Chloroplast; Thylakoid; Membrane.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-APR-1993 (Rel. 25, Created)
01-FDB-1994 (Rel. 28, Last sequence update)
28-FDB-2003 (Rel. 41, Last annotation update)
Bradykinin-potentiating peptide S4,3,2 (10C) (Angiotensin-converting enzyme inhibitor).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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15-JUL-1999 (Rel. 38, Last sequence update)
15-MAR-2004 (Rel. 43, Last annotation update)
Oxygen-evolving enhancer protein 2 (OBE2) (23 kDa subunit of oxygen evolving system of photosystem II) (Fragment).
                                                                                                                                                                                                                                                                                                                                                MEDLINE=99274068; PubMed=10344291;
Costa P., Pionneau C., Bauw G., Dubos C., Bahrman N., Kremer A.,
Frigerio J.-M., Plomion C.;
"Separation and characterization of needle and xylem maritime pine
                                                                                                                                                                            Eukaryota, Viridiplantae, Streptophyta, Embryophyta, Tracheophyta,
Spermatophyta, Coniferopsida, Coniferales, Pinaceae, Pinus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Lepidosauria, Squamata, Scleroglossa; Serpentes, Colubroidea,
Viperidae, Crotalinae, Bothrops
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             .
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80.0%; Pred. No. 3.3e+02;
tive 0; Mismatches 1; Indels
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MOD_RES 1 1 1 PYRROLIDONE CARBOXYLIC ACID.
SEQÜENCE 11 AA; 1199 MW; 20B25813C7741777 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Bothrops insularis (Island jararaca) (Queimada jararaca)
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                                                                                                                                                    inus pinaster (Maritime pine).
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Best Local Similarity
                                                                                                                                                                                                                                      NCBI TaxID=71647;
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                                                                                                                                                                                                                                                                                                                              TISSUE=Needle;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BPP3 BOTIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      proteins.
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"Primary structure and biological activity of bradykinin potentiating peptides from Bothrops insularis snake venom.";
J. Protein Chem. 9:221-227 (1990).

-I. FUNCTION: This peptide both inhibits the activity of the angiotensin-converting enzyme and enhances the action of bradykinin by inhibiting the kinases that inactivate it.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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J. Insect Physiol. 42:81-89(1996).
-I. FUNCTION: Antibacterial peptide active against Gram-negative
                                                                                                                                                                                                                                                                                                                                                                                                Bothrops insularis (Island jararaca) (Queimada jararaca).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Lepidosauria; Squamata; Scleroglossa; Serpentes; Colubroidea;
Viperidae; Crotalinae; Bothrops.
                                                                                                                                                                                                                                            01-APR-1993 (Rel. 25, Created)
01-FEB-1994 (Rel. 28, Last sequence update)
28-FEB-2003 (Rel. 41, Last annotation update)
Bradykinin-potentiating peptide 84,1,2 (Angiotensin-converting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
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Eukaryota, Metazoa, Arthropoda, Hexapoda, Insecta, Pterygota,
Neoptera, Paraneoptera, Hemiptera, Euhemiptera, Heteroptera,
Panheteroptera, Pentatomomorpha, Pentatomoidea, Pentatomidae,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 24; DB 1; Length 11; Pred. No. 3.3e+02;
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MOD_RES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1; Indels
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01-NOV-1995 (Rel. 32, Last sequence update)
28-FEB-2003 (Rel. 41, Last annotation update)
                                                                                                                                                                                            11 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TISSUE=Venom;
MEDLINE=90351557; PubMed=2386615;
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                                                                                                                                                                                      STANDARD;
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| ||||
3 GPPRP 7
                                                                                                                                                                                                                                                                                                                                                                          enzyme inhibitor)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NCBI_TaxID=55431;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Metalnikowin I
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ID BPP4 BOTIN
AC P30424;
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P80408:
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30.4%; Score 24; DB 1; Length 15;

Query Match

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TISSUE=Larva;
MEDLINE=98054539; PubMed=9392829;
Duve H., Johnsen A.H., Mestro J.-L., Scott A.G., Winstanley D., Davey M., East P.D., Thorpe A.,
"Lepidopteran peptides of the allatostatin superfamily.";
Peptides 18:1301-1309(1997).
-i- SIMILARITY: Belongs to the allatostatin family.
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Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Paraneoptera; Hemiptera; Euhemiptera; Heteroptera;
Panheteroptera; Pentatomomorpha; Pentatomoidea; Pentatomidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Cydia pomoneiia (Codiing motn).
Eukaryota, Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        30.4%; Score 24; DB 1; Length 15; 80.0%; Pred. No. 4.6e+02; tive 0; Mismatches 1; Indels
   Pred. No. 4.6e+02;
); Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Neoptera; Endopterygota; Lepidoptera; Glossata; Ditrysia;
Tortricoidea; Tortricidae; Olethreutinae; Cydia
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Antibiotic; Insect immunity.
SEQUENCE 15 AA; 1893 MW; 23835D063B946299 CRC64;
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01-NOV-1995 (Rel. 32, Last sequence update)
28-FEB-2003 (Rel. 41, Last annotation update)
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30-MAY-2000 (Rel. 39, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation update)
                                                                                                                                                      15 AA.
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                 0; Mismatches
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80.08;
                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         4; Conservative
                                                                                                                                                   STANDARD;
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 Best Local Similarity
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Best Local Similarity
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Best Local Similarity
                                           9 PRPLP 13
                                                                          8 PRPRP 12
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                                                                                                                                                                                                                                                                                                                                                           TISSUE=Hemolymph;
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                  Matches
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Best Local Similarity

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0
 Gaps
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Ascaris suum (Pig roundworm) (Ascaris lumbricoides).
Eukaryota; Metazoa; Nematoda; Chromadorea; Ascaridida; Ascaridoidea;
                                                                                                                                                                                                                                                                                                                   Cowden C., Stretton A.O.W.;
"Eight novel FMRFamide-like neuropeptides isolated from the nematode
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 01-0CT-1996 (Rel. 34, Created)
01-0CT-1996 (Rel. 34, Last sequence update)
16-0CT-201 (Rel. 40, Last annotation update)
06-0CT-201 (Rel. 40, Last annotation update)
06-year-evolving enhancer protein 2 (OEE2) (24 kDa subunit of oxygen evolving system of photosystem II) (Fragment).
Physcomitrella patens (Moss).
Bukaryota, Viridiplantae; Streptophyta; Embryophyta; Bryophyta;
NCHI_TAXID=3218;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TISSUE=Protonema;
PUDILINE=97275459; PubMed=9129336;
RABCINE=97275459; PubMed=9129336;
RABCINE=972754F, Nuske J., Reski R.;
"Çytokinin affects nuclear- and plastome-encoded energy-converting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 -!- INDUCTION: By light.
-!- SIMILARITY: Belongs to the psbP family.
Photosynthesis; Photosystem II; Chloroplast; Thylakoid; Membrane;
 0;
                                                                                                                                                                                                                                                                                                                                                            Peptides 16:491-500(1995).
-!- SIMILARITY: Belongs to the FARP (FMRFamide related peptide)
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 1; Indels
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Pred. No. 1.4e+05;
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                                                                                                                                                                                                                                                                                                                                                                                                                  9 9 AMIDATION.
9 AA; 1012 MW; 524F073774176877 CRC64;
                                                                                                                                                       01-NOV-1995 (Rel. 32, Created)
01-NOV-1995 (Rel. 32, Last sequence update)
01-FEB-1996 (Rel. 33, Last annotation update)
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100.0%; Pred. No. 100.0%; O; Mismatches
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2; Mismatches
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                                                                                                                                                                                                                                                                                                     MEDLINE=95380362; PubMed=7651904;
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Planta 201:261-272(1997).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
4; Conservative
                                                                                                                               STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                         Neuropeptide, Amidation.
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                                                                                                                                                                                                                                               Ascarididae; Ascaris.
                           1 SRAHSFG 7
                                              Query Match
Best Local Similarity
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NON TER 12
SEQUENCE 12 AA;
                                                                                                                                                                                                                                                           NCBI_TaxID=6253;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      4 PRPL 7
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                                                                                                                            FAR9 ASCSU P43172;
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P80662;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              4;
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-!- TISSUE SPECTRICITY: Brain, subosesphageal ganglion, retrocerebral complex, thoracico-abdominal ganglion, peripheral neurosecretory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota, Metazoa, Arthropoda, Hexapoda, Insecta, Pterygota,
Neoptera, Endopterygota, Diptera, Brachycera, Muscomorpha, Oestroidea,
Calliphoridae, Calliphora.
                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    "Callatostatins: neuropeptides from the blowfly Calliphora vomitoria with sequence homology to cockroach allatostatins."; Proc. Natl. Acad. Sci. U.S.A. 90:2456-2460(1993).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TISSUE=Thoracic ganglion;
MEDLINE=93211980; PubMed=8460157;
Duve H., Johnsen A.H., Scott A.G., Yu C.G., Yagi K.J., Tobe S.S.,
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57.1%; Pred. No. 5.2e+02;
tive 2; Mismatches 1; Indels
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16-OCT-2001 (Rel. 40, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
FMRFamide-like neuropeptide PF6 (NGAPQPFVRF-amide).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         system and intestine.
-!- SIMILARITY: Belongs to the allatostatin family.
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8 AA; 954 MW; D32879D5AB47740A CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                          01-NOV-1995 (Rel. 32, Last sequence update) 30-MAY-2000 (Rel. 39, Last annotation update)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Callatostatin 4 (Leu-callatostatin 4). Calliphora vomitoria (Blue blowfly).
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                                                                                                                                                                                                                                                                                                                                                                                        01-NOV-1995 (Rel. 32, Created)
                                     Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               4; Conservative
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                                                                                                                                                                                                                                                                                                                               STANDARD;
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                                                                                                                                                                   6 ANVFGAP 12
                                                                                                     3 AHSFGSP 9
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2 RPYSFG 7
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                                                                                                                                                                                                                                                                  RESULT 12
ALL4_CALVO
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FAR6_PANRE
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                        Moffett C. L. Marks N.J., Halton D.W., Thomson D.P., Geary T.G., Maule A.G.;
"Isolation, characterization and pharmacology of RMRFamide-related peptides (FaRPB) from free-living nematode, Panagrellus redivivus.";
Submitted (UUL-2000) to Swiss-Frot.
-: FUNCTION: Myoactive.
-: SIMILARITY: Belongs to the FARP (FMRFamide related peptide)
                                                                                                                                                                                                                                                      0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Purna Sai K., Jaganadham M.V., Vairamani M., Raju N.P.,
Devi A.S., Nagaraj R., Sitaram N.;
"Tigerinins: novel antimicrobial peptides from the Indian frog Rana
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hoplobatrachus tigerinus (Indian bull frog) (Rana tigerina).
Bukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi;
Amphibia; Batrachia; Anura; Neobatrachia; Ranoidea; Ranidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        tigerina.";
J. Biol. Chem. 276:2701-2707(2001).
-!- FUNCTION: Antibacterial activity against B.subtilis, E.coli,
S.aureus, M.luteus, P.putida and S.cerevisiae.
-!- SUBCELLULAR LOCATION: Secreted.
-!- TISSUB SPECIFICITY: Skin.
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16-OCT-2001 (Rel. 40, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
Pyrokinin-4 (PEP-144) (YZPL-amide).
Periplaneta fuliginosa (Smokybrown cockroach).
Bukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FUNCTION, MASS SPECTROMETRY, AND DISULFIDE BONDS.
                                                                                                                                                                                                                    Score 22; DB 1; Length 10;
Pred. No. 6.3e+02;
2; Mismatches 0; Indels
                                                                                                                                                                                                                                                   0; Indels
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                                                                                                                                                                      AMIDATION.
CB13E4C9D776C76D CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  11 AA; 1248 MW; 117D8EFD37605DCB CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                    16-0CT-2001 (Rel. 40, Created)
16-0CT-2001 (Rel. 40, Last sequence update)
10-0CT-2003 (Rel. 42, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      -1- MASS SPECTROMETRY: MW=1247; METHOD=MALDI.
Amphibian defense peptide; Antibiotic.
                                                                                                                                                                                                                                                                                                                                                                                              11 AA.
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              SEQUENCE, FUNCTION, AND AMIDATION
                                                                                                                                                                                                                    27.8%;
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                                                                                                                                                      Neuropeptide; Amidation.
                                                                                                                                                                                                                                                   3; Conservative
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PubMed=11031261;
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Best Local Similarity
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                                                                                                                                                                                                                                                                                  7 GSPRP 11
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2 GAPOP 6
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P82690;
                                                                                                                                          familv
                                                                                                                                                                                                                    Query Match
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SS DE LE
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"Tagma-specific distribution of FXPRLamides in the nervous system of the American cockroach."; J. Comp. Neurol. 419:352-363(2000).
                                                                                                                                                                                                                                                                  Gaps
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0
Neoptera; Orthopteroidea; Dictyoptera; Blattaria; Blattoidea;
Blattidae; Periplaneta.
NCBI_TaxID=36977;
                                                                                                                                                                                                                            26.6%; Score 21; DB 1; Length 12;
50.0%; Pred. No. 1.1e+03;
"Miomatches 3; Indels
                                                                                                                                                                                                        MOD_RES 12 12 AMIDATION.
SEQUENCE 12 AA; 1439 MW; F9B53049FF42CAA1 CRC64;
                                                                                                                              SEQUENCE, FUNCTION, AND MASS SPECTROMETRY
                                                                                                                                                                                                                                                                                                                                           Search completed: April 2, 2004, 10:09:17 Job time: 7.66667 secs
                                                     TISSUE=Corpora cardiaca;
MEDLINE=20189894; Pubmed=10723010;
Predel R., Eckert M.;
                                                                                                                                                                                                                                                                4; Conservative
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Best Local Similarity
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